

511/6881
FIGURE 477

TTTCCGTGCTACCTACAGAGGGGTCCATATGGCGTTGTTCTGGATTCCCGTCGTAACTCAAAGGGAAACTTTCA
CAATGTCGGGAGCCCTTGATGTCCTGCAAATGAAGGAGGAGGATGTCCTTAAGTTCCCTTGACAGCAGGAACCCACT
TAGGTGGCACCAATCTTGACTTACAGATGGAACAGTATATCTATAAAAGGAAAAGTGATGGCATCTACATCATAA
ATCTGAAGGGAACCTGGGAGAAGCTTCTGCTGGCAGCTCGTGCTATTGTTGCCATTGAAAACCTGCTGATGTCA
GTGTTATATCCTCCAGGAATACTGGCCAGAGGGCTGTGCTGAAGTTTGCTGCTGCCACTGGAGCCACTCCAATTG
CTGGCCACTTCACTCCTGGAACCTTCACTAACCAGATCCAGGCAGCCTTCTGGGAGCCATGGCTTCTTATGGTTA
CTGACCCCAGGGGCTGACCATCAGCCTCTCACAGAGGCATCTTATGTTAATCTACCTACCATCGCTCTGTGTAAC
ACAGATTCTCCTCTGCGCTATGTGGACATTGCCATCCCGTGCAACAACAAGGGAGCTCACTGAGTGGGTTTGATG
TGGTGGATGCTGGCTCGGGAAGTACTGTGCATGCGTGGCACCATTTCCCGTGAACACCCATGGGAGGTCATGCCT
GATCTCTGCTTCTACAGAGATCCTGAAGAGATTGAAAAAGAAGAGCAGGCTGCTGCTGAAAAGGCAGTGACCAAG
GAGGAATTTAGGGTGAATGGTCTGCTCCAGCTCCTGAGTTCACTGCTACTCAGCCTGAGGTTGCAGACTGGTTT
GAAGGTATACAGGTGCCCTCTGTGCCTATTACAGCAGTTCCTACTGAACAATGGAGCACTCAGCCTGCCACGGAAT
ACTGGTCTGCAGCTCCCACTGCTCAGGCCACTGAATGGGTAGGAGCAACCACTGAATGGTCTTAAGCTGTTCTTG
CATGGGCTCTTAAGCAACATGGAAAAATGGTTGATGGAAAATAAACATCAGTTTCT

512/6881
FIGURE 478

MSGALDVLQMKEEDVLKFLAAGTHLGGTNLDLQMEQYIYKRKSDGIYIINLKGTWEKLLLAARAIVAIENPADVS
VISSRNTGQRAVLKFAAATGATPIAGHFTPGTFTNQIAAFWEPWLLMVTDPGR

513/6881
FIGURE 479

AAAGGGAACTTTTACAATGTCCGGAGCCCTTGATGTCCTGCAAATGAAGGAGGAGGATGTCCTTAAGTTCCTTG
CAGCAGGAACCCACTTAGGTGGCACCAATCTTGACTTACAGATGGAACAGTATATCTATAAAAGGAAAAGTGATG
GCATCTACATCATAAATCTGAAGGGAACCTGGGAGAAGCTTCTGCTGGCAGCTCGTGCTATTGTTGCCATTGAAA
ACCCTGCTGATGTCAGTGTTATATCCTCCAGGAATACTGGCCAGAGGGCTGTGCTGAAGTTTGCTGCTGCCACTG
GAGCCACTCCAATTGCTGGCCACTTCACTCCTGGAACCTTCACTAACCAGATCCAGGCATCTTATGTTAATCTAC
CTACCATCGCTCTGTGTAAACACAGATTCTCCTCTGCGCTATGTGGACATTGCCATCCCGTGCAACAACAAGGGAG
CTCACTGAGTGGGTTTGATGTGGTGGATGCTGGCTCGGGAAGTACTGTGCATGCGTGGCACCATTTCCTGTAAC
ACCCATGGGAGGTCATGCCTGATCTCTGCTTCTACAGAGATCCTGAAGAGATTGAAAAAGAAGAGCAGGCTGCTG
CTGA

514/6881
FIGURE 480

MSGALDVLQMKEEDVLKFLTAGTHLGGTNLDFQMEQYIYKRRSDGIYIINLKRTWEKLLLAARAIVAIENPAVVS
VISSRNTGQRAVLKFAAVTGATPIAGRFTPGNFTNQIQEASYVNLPTIALCNTDSPLHFVDIAIPCNNKGAH

515/6881
FIGURE 481A

AGGTAGCAAGATGGCCGCCGCTGAGGAAGGCTGTAGTGTGCGGGCCGAAGCGGACAGGGAATTGGAGGAGCTTCT
GGAAAGTGCTCTTGATGATTTTCGATAAAGCCAAACCCCTCCCCAGCACCCCTTCTACCAACACGGCCCTGATGC
TTCGGGGCCCCAGAAGAGATCGCCAGGAGACACTGCCAAAGATGCCCTCTTCGCTTCCCAAGAGAAGTTTTC
GGAACCTATTTCGACAGTGAACCTGGCTTCCCAAGCCACTGCGGAGTTCGAGAAGGCAATGAAGGAGTTGGCTGAGGA
AGAACCCACCTGGTGGAGCAGTTCACAAAGCTCTCAGAGGCTGCAGGGAGAGTGGGCAGTGATATGACCTCCCA
ACAAGAATTCACTTCTTGCCCTAAAGGAAACACTAAGTGGATTAGCCAAAAATGCCACTGACCTTCAGAACTCCAG
CATGTGCGGAAGAAGAGCTGACCAAGGCCATGGAGGGGCTAGGCATGGACGAAGGGGATGGGGAAGGGAACATCCT
CCCCATCATGCAGAGTATTATGCAGAACCTACTCTCCAAGGATGTGCTGTACCCATCACTGAAGGAGATCACAGA
AAAGTATCCAGAAATGGTTGCAGAGTCATCGGGAATCTCTACCTCCAGAGCAGTTTGAAAAATATCAGGAGCAGCA
CAGCGTCATGTGCAAAATATGTGAGCAGTTTGAGGCAGAGACCCCCACAGACAGTGAACCACTCAAAAGGCTCG
TTTTGAGATGGTGTGGATCTTATGCAGCAGCTACAAGATTTAGGCCATCCTCCAAAGAGCTGGCTGGAGAGAT
GCCTCCTGGCCTCACTTTGACCTGGATGCCCTCAATCTTTCGGGGCCACAGGTGCCAGTGGTGAACAGTGTCT
GATCATGTGAACACAACACGTTTTCTCTCTGAGTCCCAGCTATGGGGAACATCTGGAGTCAGCAGAACCATTG
GGACCTGAGGCAGGAGTGTACCTGCGGGAGAAGTCTGCCCGCTGCCCTCTGTATCCCATTCAGATTGTGCCA
TACCAGCTGAGGTTTTCTCTGTCTCTCTAGGAATAGGGTCTGTTTTCACAGGCCATTTCTGTGAACCTACTCC
ATTGTGGTTTTCTGCCACTATCAAAGTTCCAGCTACCTGCAAGGTGAAGGAAGGCATCCCTTTTGGGGCATGCACT
TTCTTTCCCTTTCTCAAAATAATGTTATATGTGGCCACACTGATGTTACCTTTACGTCCAGGGTCTTTGTGCCCT
GTCTCTACTCCCTCTCTTGGATCTGGGGAGGAGGGGCAGAGACCTGGGACTCTGTATTTCTATAGTTCTCTCTGGC
AGAGCCCTTTGAGAATGGGGAGAAACAGCCTGGGCTGGGGCTACAGGTCTGTCACTATGTCTCTTGCCTTCAGAC
AGACCATTCTGAATTTCTTAAAGGGAAGGGCTTTTGCATCTAATCACAATAGAGTTGAAAGAGAGGCCTTAGGA
TTCTCTCTCTCTAGGTGTGAGCCCTCACCTCCCTGTTCCAGGCTGAGAACTCAAATGGTTACCCTGCTTCTTC
CTACAATGCTGTGTGATATGGGTGAACCCAGCCCCCTGACCTTCTCTATCCCTGCCCCATCTCCCTTTTACCTC
CTCTCTTTTTTAAACACCTGTTTATCCCAACCTTTTTGAGCTCAAGCTGTGATAAAGAAGGGCCCATCTATTTT
CCCTCATCTAGTCCATTTACGATTCTCACTGACTCCCCGTCTCTCTGGCAGACACAAATAAACCAGTGTCAAGT
CTAGGAAATTAATGGCTATTCTTCCCCAGATACATTCTGGCTTATTTGAGATACATGATTCTCTTAGAATCCTGT
CCCTTGGTTTCAAGGAAAGTAGCTTGGAAAAGGAGTAGGGGTATAGCTTGGGTCCCTTTTCTGCAAGGCCCATGG
GGCAGAAATATAATAATATTCTGAGTGAGGAGTGTGGTCTTTTTCTGATCTTCTCAGCTTCCGTAAGTTGCAGA
GTGAGGTATATTAGGAGACTAGTTCTACACAATATTGTAATGTCTGGGTCCATCAACACCCACCTTCCACAACCT
AGTCTGCACCTCAGTTGGCAAAGGAGACTGGATGGCCATCTTCTCTCATGTTCCCTTGAGTATTTCAATGTAGAA
AGCCCTTCAAGTGGTATTATATTTTAAACCTTTTACATTATTTGTTATTAATGTTAGTAATATATTGTTATGTTTT
TAAATTATTTTTCTTTAAGCTGACGTGGCTTTTTTCTGTGGCTCCAGTGGGTCTACGGACCTTGGCTGACATA
TGTTGGTAGGTACTCTGGTCAGCTCAGCTGGCTGTCTGGTTCCTCAGACAGATAAGTCTCTCCAAGCAAACTTC
ACATGCATTATGAGTCGCTTTGAGCTTCTGACATGTCACTTGGCCCGAGGTTAAACTTTTACCCTTTGAAGAC
CTTACATGTTTTATGGTATTGGTGAGGAAGGAAATGTTCTCAAGGTCTCAGGCTATTGGGAAATTCCAACCTCCT
ATACCTTACCAGAGCATGGAAGAGCCCAGATCTGAATGTAAACGTCTCTGTTCTGCCAGAGATGGAAAAAATAC
AGGTATACTTGTGATATAGTCATGGGGCTTCAGTGTCACTATTTCTCTCTTAAAGCTCCAGCCAAAACTGGACA
AGGATAGAGAGGAGGAGGGAAGAACAAAAGAGCCCTTCTCTATGAACCTTGTGCCTTCTGTCTTACCAGTTTTCT
TTTACAGATTCTCACTTCTGCTAGCCTAGCCAGGGCTTACTCCAGGAATCTAAATAGATGCCCTAGTCCACTTTA
TCTTTGTTCCCAAGGCACTCATTTTTATTTTGATTTGATTGAATGTGAGCAGGTGACCTCAGGTACACTTTG
TTCCAAAACTTTTGAATTATTCAGGACTTGTGGTGGAGTTATGGTACTCTAGGGCAGTCTTTCTCAAACTAT
GTATGGTAAAGGACAGGTTTTTGTTTTCCAGTCCCTCACTTATCAATATGCATTCTTATTGCCGATGACAGGT
ATGGAGTTCACACTGTGTGCTGCCGACCCGGCAAGTTTGACAGCACCCAACTGGCCAGACTGTTCTGTAGGTTA
AGTCCATTGTATCATGTACTTGATATCACAGCAACATTGAAATGTAAAAAGTTTTTAAACACTCTCAATTTCTA
ATTACCATGTACAGACTGGTGAAAAAAGGTTGTTCACTGACCAGCACAAAGTCTGCAGATCATCTTTG
AGTAGCACTGTTTTGGGGCCCTCGGTCTCTCTGAAGACCCTAGCAGAACTGATACCTACCTGTATCTCTTGTCT
CTCCTATTTGAGTTTCACTTCCAGAGAACTTGTCTTCAAGCAAGAAATGTGTCACTAGTAAGGACATCTCTAGCAT
TTCTCTAGCCTTCTTTTTCTGCTGCTCAAAATAATCGTTACAAAGCTTAGGTTTAAAGCTGTATATGAAATATTT
ATGCGACTCTCAAACTTTAAAGGAGTTGCTCCTTTGTTCCAAATTAATGTGTTAGATAAAATTTGTGATTGTAT

516/6881
FIGURE 481B

GGGTGGCTTCATGAATTAAGAATTGAATTAATACAGACTTTTTGATAATAAAAAAAAAAAAAA

517/6881
FIGURE 482

MAAAEEGCSVGAEADRELEELLESALDDFDKAKPSPAPPSTTTAPDASGPQKRSPGDTAKDALFASQEKFFQELF
DSELASQATAEFEKAMKELAE EEPHLVEQFQKLSEAAGRVS DMTSQQEFTSCLKETLSGLAKNATDLQNSSMSE
EELTKAMEGLGMDDEGDGEGNILPIMQSIMQNLLSKDVLPSLKEITEKYPEWLQSHRESLPPEQFEKYQE QHSVM
CKICEQFEAETPTDSETTQKARFEMVLDLMQQLQDLGHPPELAGE MPPGLNFDLDALNLSGPPGASGEQCLIM

518/6881
FIGURE 483A

GAGAAGGGACCTTCAGGTCCAGGCAAAGGGGGAACCTTCTGTCGTGGGAACGAAAAAGAAAGAGGATTTACAGGGT
GGGGGGACAGAGGGGCGAGCAGGAACCAGAAGGGAGACAGTGGCGGTGCGGGCGGGGCCGATCCGAGAGTTCCCT
TAGAGAACGGAGCTCACGGGCGGGGAGGCCTCACCTGCTAGTAGGACGCAGAAAGACAGAAGGCCAAGGAGACCC
CGACTTCCCGGGTCAGCCCCAGAGCCACCCCTGCCGTAGCCATCTTGCTCTCTGCTGAGCGGAAGCCCCCGTT
CGGCTCCTGTCTGTTAGCGGCTCTCTAGGCTACCACTGACACCGTCTCTGTGGCCCGGAGCCTAAGAGACCGGA
AGTTCGTGTTTCCAGGCGCTTCCGGAAACCGCGGGAGAGGGTCGCTGACGTGGAGGCGTCCGAAGGGCAGCAGGG
TGTGTCGGGGCTCGGATTAAAGACATCGGAGTCGGAGACCTGAGAGATGTTAACCAAATTCGAGACCAAGAGCGCG
CGGGTCAAAGGCTCAGCTTTCACCCCAAAAGACCTTGGATCCTGACTAGTTTACATAATGGGGTCATCCAGTTA
TGGGACTATCGGATGTGCACTCTCATTGACAAGTTTGATGAACATGATGGTCCAGTGGCAGGCATTGACTTCCAT
AAGCAGCAGCCACTGTTCTGCTCTGAGGAGATGACTATAAGATTAAGGTTTGGAAATTACAAGCTTCGGCGCTGT
CTTTTCACATTGCTTGGGCACTTAGATTATATTCGCACCACGTTTTTTCATCATGAATATCCCTGGATTCTGAGT
GCCTCCGATGATCAGACCATCCGAGTGTGGAACGGCAATCTAGAACCTGTGTTTGTGTGTTAACAGGSCACAAC
CATTATGTGATGTGTGCTCAGTTCACCCACAGAAAGACTTGGTAGTATCAGCCAGCCTGGACCAGACTGTGCGC
GTTTGGGATATTTCTGGTCTGAGGAAAAAAACCTGTCCCTGGTGGGTTGGAATCGGATGTGAGAGGAATAACT
GGGGTTGATCTATTTGGAACCTACAGATGCAGTGGTGAAGCATGTACTAGAGGGTCACGATCGTGGAGTAACTGG
GCTGCCTTCCACCCCACTATGCCCTTATTGTAICTGGGGCAGATGATCGTCAAGTGAAGATCTGGCGCATGAAT
GAATCAAAGGCATGGGAGGTTGATACCTGCCGGGGCCATTACAACATGTATCTTGTGCCGTCTTCCACCTCGC
CAAGAGTTGATCCTCAGCAATTCTGAGGACAAGATATTGAGTCTGGGATATGTCTAAGCGGACTGGGGTTCAG
ACTTTCGCAGAGACCATGATCGTTTCTGGGTCTAGCTGCTCACCTAACCTTAACCTCTTGCAGCAGGCCAT
GATGGTGGTATGATTGTGTTTAAAGCTGGAACGGGAACGGCCAGCCTATGCTGTTTATGGCAATATGCTACACTAT
GTCAAGGACCGATTCTTACGACAGCTGGATTTCAACAGCTCCAAAGATGTAGCTGTGATGCAGTTGCGGAGTGGT
TCCAAGTTTCCAGTATTCAATATGTACATAATCCAGCAGAAAAATGCAGTCCCTGCTTTGTACAAGAGCTAGCAAT
CTAGAGAATAGTACCTATGACCTGTACACCATCCCTAAAGATGCTGACTCCCAGAATCCTGATGCGCCTGAAGGG
AAACGATCCTCAGGCCTGACAGCCGTTTGGGTGCTCGAAATCGGTTTGTGTCCTAGATCGGATGCATTTCGCTT
CTGATCAAGAATCTGAAGAATGAGATCACCAAAAAGGTACAGGTGCCCACTGTGATGAGATCTTCTATGCTGGC
ACAGGCAATCTCCTGCTTCGAGATGCGGACTCTATCACACTCTTTGACGTACAGCAGAAGCGGACTCTGGCATCT
GTGAAGATTTCTAAAGTGAAATACGTTATCTGGTCAGCAGACATGTACATGTAGCACTACTAGCCAAACACGCC
ATTGTGATCTGTAAACGCAACTGGATGCTTTATGTAACATTATGAGAACATTCGTGTCAAGAGTGGGGCCTGG
GATGAGAGTGGGGTATTATCTATACCAAGCAACCACATCAAAATATGCTGTCAACACTGGGGACCACGGGATC
ATTCGAACCTCTGGATTTACCCATCTATGTACACGGGTGAAGGGCAACAATGTATACTGCCTAGACAGGGAGTGT
CGTCCCCGGGTACTACCATTTGATCCCACTGAGTTCAAATTCAGAGCTGGCCCTGATCAACAGAAAATATGATGAG
GTACTGCACATGGTGAGGAATGCCAAACTAGTTGGCCAGTCTATTATTGCTTATCTCCAGAAGAAGGGCTATCCT
GAAGTGGCACTGCATTTTGTCAAGGATGAGAAAACCTCGCTTAGTCTGGCACTGGAGTGTGGAACATTGAGATT
GCTCTGGAAGCAGCCAAAGCACTGGATGACAAGAACCTGCTGGGAAAAGCTGGGAGAAGTGGCCCTGCTGCAGGGG
AACCACCAGATTGTGGAATGTGCTATCAGCGTACCAAAAACCTTTGACAACTTTCTTCTGTATCTTATCACT
GGCAACTTAGAAAAACTTCGCAAGATGATGAAGATTGCTGAGATCAGAAAGGACATGAGTGGCCACTATCAGAAT
GCCCTATACCTGGGTGATGTGTGAGAGCGTGTGCGGATCCTGAAGAACTGTGGACAGAAGTCCCTGGCCTATCTC
ACAGCTGTACCCATGGCTTAGATGAAGAAGCTGAGAGCCTAAAGGAGACATTTGACCCAGAGAAGGAGACAATC
CCAGACATTGACCCTAATGCCAAGCTGCTCCAGCCACCTGCACCTATCATGCCATTGGATACCAATTGGCCTTTA
TTGACTGTATCCAAAGGATTTTTTGAAGGCACCATTTGCCAGCAAAGGGAAGGGAGGAGCACTGGCTGCTGACATT
GACATTGACACTTTGGTACAGAGGGCTGGGGAGAGGATGCAGAGCTGCAGTTGGATGAAGATGGGTTTGTGGAG
GCTACAGAAGGTTTGGGGGATGATGCTCTTGGCAAGGGACAGGAAGAAGGAGGTGGCTGGGATGTAGAAGAAGAT
CTGGAGTCCCTCCTGAGCTGGATATATCCCTGGGGCAGCTGGTGGGGCTGAAGATGGTTTCTTTGTGCCCCCA
ACCAAGGGAACAAGTCCAACCTCAGATCTGGTGAATAACTCTCAGCTTCCAGTTGATCACATCTGGCAGGCTCT
TTCGAAACAGCCATGCGGCTCCTTCATGACCAAGTAGGGGTAATCCAGTTTGGCCCTACAAGCAACTGTTCTTA
CAGACATACGCCCCGAGGCCGACAACCTATCAGGCTCTGCCCTGCCTACCCTCCATGTATGGCTATCCTAATCGC
AACTGGAAGGATGCAGGGCTGAAGAATGGTGTACCAGCTGTGGGCTGAAGCTTAATGACCTCATCCAACGGTTG
CAGCTGTGCTACCAGCTCACCACAGTTGGCAAATTTGAGGAGGCTGTGGAAAAATTCGGTTCCATCCTTCTCAGT

519/6881
FIGURE 483B

GTGCCACTTCTTGTGTGGACAATAAACAAGAGATTGCAGAGGCCAGCAGCTCATCACCATTGCGGTGAGTAC
ATTGTGGGTTTTGTCCGTGGAGACAGAAAGGAAGAAGCTGCCCAAAGAGACTCTAGAACAGCAGAAGCGCATCTGT
GAGATGGCAGCCTATTTACCCACTCAAACCTGCAGCCTGTGCACATGATCCTGGTGCTGCGTACAGCCCTCAAT
CTGTTCTTCAAGCTCAAGAACTTCAAGACAGCTGCCACCTTTGCTCGGCGCCTACTAGAACTCGGGCCCAAGCCT
GAGGTGGCCCAACAGACCCGAAAAATCCTGTCTGCCTGTGAGAAGAATCCACAGATGCCTACCAGCTCAATTAT
GACATGCACAACCCCTTTGACATTTGTGCTGCATCATATCGGCCCATCTACCGTGGAAGCCAGTAGAAAAGTGT
CCACTCAGTGGGGCTGCTATTTCCCTGAGTTCAAAGGTCAAATCTGCAGGGTCACCACAGTGACAGAGATTGGC
AAAGATGTGATTGGTTTAAGGATCAGTCTCTGCAGTTTCGCTAAGGCCCTTTGTGTGCATGGGTGAGTCACC
ATATGTTCCCCCAGAGAATGTGTCTATATCCTCTTCTAACAGCACCTTCCCCCTGCAGTACTCTTCAGATCT
GGCTCTCTGTACCCTAAAACCTAGTATCTTTTTCTCTTCTATGGAATCCGAAGGTCTAAACTTGACTTTTTTG
AGGTCTTCTCAACTTGACTACAGTTGTGCTCATAATTGTCCTTGCCCTTCCAGCTTAATTATTTAAGGAACAAA
TGAAACTCTGGGCTGGGTGGAGTGGCTCATACCTGTAATCCCAGCACTTTGGGAGGCTACGGTGGGCAGATCAT
CTGAGGCCAGGAGTTCGAGACCTGCCTGGCCAACATGGCAACACCCCGTCTCTAATAAAAAATATAAAAAATTAGCC
TGGCATGGTAGCATGCGCTATAGTCCCAGCTGCTCAGGAGGCTGAGGCATGAGAATCGCTTGAACCTAGGAGGT
GGAGGTGCAATTCAACTGAGATCATACCACTTCATTCCAGCCTGGGTGACAGAGCAAGACTCTGTCTCAAAAAA
AAAAAAGGAAACTCTGTGATGGACATTTGTTTAGTAAATCCCTTCAGTATTTATCCCTCCTTTCCCCACAGCA
GCTTCTTCTCTGTCAACTAGAAAGGAGCAGGATGTAATAAATACATTTTGGTGTGACTAGGCCACACCAACTCT
TAATCATCTCCCATTTTCTTAGACATTTAAATTTCAAGGCAGGTACCCTCTGTGTACTCAGAAATTTGAAGAAG
TTATTTGGTTTTCCAAATGCACACTGCGGGTTATTGATTTGTTCTTTACAACATATTGTTCTCATATTTCTCACA
CTAAATAAATCTCTATGAGAGCTTCTTG

520/6881
FIGURE 484

AGCTGCGGTGTTGTGCTGTGGGAAGGGAGAAGGATTTGTAAACCCCGGAGTGAGGTTCTGCTTACCCGAGGCCG
CTGCTGTGCGGAGACCCCGGGTGAAGCCACTGTCATCATGTCTGACCAGGAGGCAAACCTTCAACTGAGGACT
TGGGGGATAAGAAGGAAGGTGAATATATTAACTCAAAGTCATTGGACAGGATAGCAGTGAGATTCACTTCAAAG
TGGTTCCAATGAATTCATTCAGGTTTCTTTGAGGGTCAGAGAATTGCTGATAATCATACTCC

521/6881
FIGURE 485

GAAGGATTTGTAAACCCCGGAGTGAGGTTCTGCTTACCCGAGGCCGCTGCTGTGCGGAGACCCCGGGTGAAGCC
ACTGTCATCATGTCTGACCAGGAGGCAAAACCTTCAACTGAGGACTTGGGGGATAAGAAGGAAGGTGAATATATT
AAACTCAAAGTCATTGGACAGGATAGCAGTGAGATTCATTCAAAGTGAAAATGACAACACATCTCAAGAACTC
AAAGAATCATACTGTCAAAGACAGGGCGTTCCAATGAATTCATTCAAGTTTCTCTTTGAGGGTCAGAGAATTGCT
GATAATCATACTCCAAAAGAACTGGGAATGGAGGAAGAAGATGTGATTGAAGTTTATCAGGAACAAATGGGGGGT
CATTCAACAGTTTAGATATTCTTTTTATTTTTTTCTTTCCCTCAATCCTTTCTCATTTTTAAAAATAGTTCTT
TTGTAATGTGGTGTTAAAAACGGAATTGAAAACCTGGCACCCCATCTCTTTGAAACATCTGGTAATCTGAATTCTA
GTGCTCATTATTATTGTTTATTTTCATTGTGCTGATTTTGGTGATCAAGCCTCAGTCCCCTTCATATTAC
CCTCTCCTTTTTATTGTGGTGATAAATAAGATTGACCAATGCAAGCGTTCATAATAACTTGCGAATTGGCCCTGA
TGTTCTAGCATGTGATTACTTCACT

522/6881
FIGURE 486

GTTTGCCAAAAATCCCAGGCAGCATGGACCTCAGTCTTCTCTGGGTACTTCTGCCCTAGTCACCATGGCCTGGGG
CCAGTATGGCGATTATGGATACCCATACCAGCAGTATCATGACTACAGCGATGATGGGTGGGTGAATTTGAACCG
GCAAGGCTTCAGCTACCACTGTCCCCAGGGGCAGGTGATAGTGGCCGTGAGGAGCATCTTCAGCAAGAAGGAAGG
TTCTGACAGACAATGGAACCTACGCCTGCATGCCACGCCACAGAGCCTCGGGGAACCCACGGAGTGCTGGTGGGA
GGAGATCAACAGGGCTGGCATGGAATGGTACCAGACGTGCTCCAACAATGGGCTGGTGGCAGGATTCCAGAGCCG
CTACTTCGAGTCAGTGCTGGATCGGGAGTGGCAGTTTTACTGTTGTCGCTACAGCAAGAGGTGCCCATATTCTTG
CTGGCTAACAACAGAATATCCAGGTCATATGGTGAGGAAATGGACATGATTTCTACAATTATGATTACTATAT
CCGAGGAGCAACAACCACTTTCTCTGCAGTGGAAAGGGATCGCCAGTGGAAGTTCATAATGTGCCGGATGACTGA
ATACGACTGTGAATTTGCAAATGTTTAGATTTGCCACATACCAAATCTGGGTGAAAGGAAAGGGGCCGGGACAG
GAGGGTGTCCACATATGTTAACATCAGTTGGATCTCCTATAGAAGTTTCTGCTGCTCTCTTTCTCTCCCTGAG
CTGGTAACTGCAATGCCAACTTCCTGGGCCTTTCTGACTAGTATCACACTTCTAATAAAATCCACAATTAACCA
TGTTTCTCACTTTTACATGTTTCATAGCAACTGCTTTATATGACTGATGATGGCTTCCTTGCACACCACATATA
CAGTGCGCATGCTTACAGCCGGGCTTCTGGAGCACCAGCTGCAGCCTGGCTACTGCTTTTACTGCAGAATGAAC
TGCAAGTTCAGCATAGTGGAGGGGAGAGGCAGAACTGGAGGAGAGGTGCAGTGAAGGTCTCTACAGCTAAGCCT
GTTTGAATGATACGTAGGTTCCCCACAAAAGCAGGCTTTCTGCCCTGAGGGACATCTTCCCACTCCCTGCTCC
ACATGAGCCATGCATGCTTAGCAATCCAAGTGCAAGAGCTCTTTGCTCCAGGAGTGAGGAGACTGGGAGGTGAAAT
GGGGAAATGGAAGGGTTTGGAGGCAGAGCTGAAAACAGGGTTGGAAGGATTTCTGAATTAGAAGACAAACGTTA
GCATACCCAGTAAGGAAAATGAGTGCAGGGGCCAGGGGAACCCGTGAGGATCACTCTCAAATGAGATTAAAAACA
AGGAAGCAGAGAATGGTCAGAGAATGGGATTAGATTGGGAACCTGTGGGGATGAGAGTGACCAGGTTGAACTGG
GAAGTGAAAAAGGAGTTTGTAGTCACTGGCACCTAGAAGCCTGCCCACGATTCTTAGGAAGGCTGGCAGACACC
TGGAACCCTGGGGAGCTACTGGCAAACCTCTCTGGATTGGGCCTGATTTTTTTGGTGGGAAAGGCTGCCCTGGGG
ATCAACTTTCCTTCTGTGTGTGGCTCAGGAGTTCTTCTGCAGAGATGGCGCTATCTTTCCTCCTCCTGTGATGTC
CTGCTCCCAACCATTTGTACTCTTCATTACAAAAGAAATAAAAAATATTAACGTTT

523/6881

FIGURE 487

MDLSLLWVLLPLVTMAWGQYGDYGYQYHDYSDDGWVNLNRQGFQCPQGQVIVAVRSIFSKEGSDRQWNY
ACMPTPQSLGEPTECWEEINRAGMEWYQTCSNGLVAGFQSRIFESVLDREWQFYCCRYSKRCPYSCWLTTEYP
GHYGEEMDISYNDYYIRGATTTFAVERDRQWKFIMCRMTEYDCEFANV

524/6881
FIGURE 488

AGAGGAAGCCAAGAACGGTGAAAAGGCCAGGCGGAGTTCAGAGGAGGTGGACGGCCAGCACCCGGCCCCAAGAGGA
GGTCCCGGAATCGCCCCAGACCTCTGGCCCAGAGGCAGAAAAATAGGTGTGGGAGCCCCAGGGAGGAAAAGCCAGC
TGGAGAGGAAGCAGAGATGGAAAAAGGCTACAGAGGTGAAGGGGGAGAGGGTGCAAAATGAAGAGGTGGGACCTGA
ACATGACAGCCAAGAAACAAGAAGCTGGAGGAGGGAGCTGCAGTGAAGGAGACCCCCACAGTCCCCCTGGAGG
AGTGAAGGGCGGAGATGTCCCAAGCAGGAAAAAGGCAAGGAAAAACAACAGGAGGGGGCAGTGTCTGAGCCAGG
CTGCAGCCCCCAGACCGGCCCTGCCAGCTGGAGACCAGCAGTGAGGTCCAGAGCGAGCCAGCAGTCCCCAAGCC
GGAGGATGACACTCCTGTCCAGGACACTAAAATGTGAAGAACAGCTCATTGTGCCCCAGTGATGAAGTTGCTGGA
CACATCTCTTTGCAGGTAGCAGCAACAGTTGTAGCAGCAGCAGACGAAGCCATTGCAGAGGCAGAAATATGCTGAG
TGTCTGGAGTCAGCCTGAAGACACAGGGTGGATTATTTCTGGCCTCCACACCAAACGTTCCCTTGCAGATGGAG
ACTGAATCTGAGGGCAGCAGACTTTTATCAGCTTGAGTTTATGTCATTTGATGGACTTGGTTCAACAACAAGAAC
TTACTTAAACAATGTACTGTGGTGATGAGTCCAGGGGCACTGGTCAGCCTGTGGAGCCCTGGATGCTATCCAC
ACCCACCTATCCCTGCAGCTAATTTAGCTGATCTCTAATTTAACTGAGCTCTAATTTAGCTGATCAGATTTTGCT
TGGGTAAAGTTCCTTTTTAATGTTCTAAAGTGTTCACGGTTCTCAAATATCAGTTAAAACTAATTTTAGGTGGC
CATAAACATAAAATAGAAACCCCTGTAAGTTACAGAAGACCCTAAATTGTATCAAAACCCCTAGAGACAACCTTTCA
ATTTGATCCAAATTTGAAGTGGCCAACAGTCTTTAAACACTGGACTAGAAGAGATAATGATTGAAACATTTAA
AAAAAAAAGTGTCCATTTCGAGGAGCTTTTCTGTCTGTGGTTTCCAGTTGGTGACCACCATGGGAGGTCTG
CTGGCTCGGCTCACTCCCTTCTCCACCCCTTGAGAATGTGGAGAACTCCCATGGAGAGGCAGAAATGGCAGGAGGT
TTCATGTCCCGGTTGCATCTCTCTCTGAAAGAAAAGCAGTGATACCTGAATAATGCTGGCTCTCCGATTGATCC
TGTGAGGATGAATTTGCATTTCCAGAATCCTTGAGCATGGATTAGATGTTTCTGGGAGGTGCCTTGAGTACCAT
TATGTGCAAGCTACATAATTTAAACATTTTTCTTAGTTTTCCCTGGGAAGCTTTTCTTGACTCACAGCCCAGGTTC
TTCTGCCCAACACAAAAGGAGTGAGTTGGGGTCTTTAGTCTCTTCTTATTGGGTAGCTCTTGCTTTAATATCTG
TTTGGTGAGTGTAAGGGATTCTGCAAGGGACAGGGGGCTGACTACCCAGTCTTTGACTTGATCCTCTCCCCTC
TTCATACACTCCTGCTGAAAAATGTTAATCCAAATACACATTTAACTTAGGGTCGGTCCTTATTCTGATTTGAG
TATTTTAATGTCTCAGTGTGCTGATTTGGTAGTTGGAAGAATTATTCTTCTGGAGGTCTGTTAGACTACATCCTA
CACTGACTTCAGAAAACAGTCTGTGAGACAAAAGGCCTTATGTCAACCACTGGTACCTCAGTTTCCTCATCCCAT
TTACAGTTTTTCTAACTCCAGGGTAGTGTTTAGTGTTAATATTTGGGATATATTTTTTTTCAAACTGTTTTTAA
GTAGTTTGTAATTTGTAACAACTTGTAACCTGGTTGGGACTGATATTGTCATAGCTATGATAAACTTTGGATAT
TAGCAGAATTTGGGAAAAAAAAAAAAAAAAAAAAA

525/6881
FIGURE 489

MEKATEVKGERVQNEEVGPEHDSQETKKLEEGAAVKETPHSPPGGVKGGDVPKQEKGEKQQEGAVLEPGCSPQT
GPAQLETSSEVQSEPAVPKPEDDTPVQDTKM

526/6881
FIGURE 490

GGACGTTGAGAGAACGAGGAGGAAGGAGAGAAAATGGCATCCATGGATTACAGTACCTATAGCCAAGCTGCAGCG
CAGCAGGGGCTACAGTGCTTACAACACCCAGCCCACTCAAGGATATGCACAGACCACCCAGGCATATGGGCAACAA
AGCTATGGAACCTATGGACAGCCCATTTGATGTCAGCTATACCCAGGCTCAGACCCTGCAATCTATGGGCAGACC
GCCTATGCAACTTCTTATGGACAGCCTCCCCTGTTTACTACTCCAAGTGGGGGAGGCATACAGCCAGCCT
GTCCAGGGGTATGGCACTGGTGCTTATGATACCACTGCTACAGTCACCACCACCCAGGCCTCCTATGCAGCT
CAATCTGCATATGGCACTCAGCCTGCTTATCCAGCTTATGGGCAGCAGCCAGTAGCCACTGCACCTACAAGACTG
CAGAATGGAACAAGCCCACTGAGACTAGTCAACCTCAATCTAGCACAGGGGGTTACAACCAGCCAGCCTAGGA
TATGGACAGAGTAACTGCAGTTATCCCCAGGTACCTGGGAGCTACTCCATGCAGCCAGTCACCGCACCTCCATCC
TACCCTCCTACCAGCTATTCTCTACACAGCCAAGTAGTTATGATCAGAGCAGTTACTCTCAGCAGAACACCTAT
GGGAAACCGAGCAGCTATGGACAGCAGAGTAGCTATGGTCAACAAAGCAGCTATGGGCAGCTGCCTCCCCTAGT
TACCCACCCCAAACTTGATCCTACAGCCAAGCTCCAAGTCAGTATAGCTAACAGAGCAGCAGCTACGGGCAGCAG
AGTTCATTCTGACAGGACCACCCAGTAGCATGGGTGTTTATGGGCAGGAGTCTGGAGGATTTTCCGGACTAGGA
GAGAACCGGAGCATGAGTGGCCCTGATAACTGGGGCAGGGGAAGAGGGGGATTGATCGTGAGGCATGAGCAGA
GGTGGGCGGGGAGGAGGATGCGGTGGAATGGGCAGCGCTGGAGAGCAAGTTGGCTTCAATAAGCCTGGTGGACCC
ATGGATGAAGGACCAGATCTTGATCTAGGCCCACCTGTAGATCCAGATGAAGACTCTGACAACAGTGCAATTTAT
GTACAAGGATTAAATGACAATGTGACTCTAGATGATCTGGTAGACTTCTTTAAGCAGTGTTGGGTTGTTAAGATG
AACAAGAGAAGTGAAGCAACCCATGATCCACACCTACCTGGACAAGGAAACAAGAAAGCCCAAGGTGATGCCACA
GTGTCCTGTGAAGACTCACCTACTGCCAAAGCTGCCGTGGAATGGTTTGATGGGAAAGATTTCAAGGGAGCAAA
CTTAAAGTCTCTCTTGCTCGGAAGAGGCCTCCAGTGAACAGTATGCAGGGTGGTATGCCACCCCATGAGGGCAGA
GGGATGCCACCACCACTCTGCGGAGGTCCAGGAGGCCAGGAAGTCTGGGGGACCCATGGGTACATGGGAGGC
CGTGGAGGAGATAGAGGAGGCCTCCCTCCAAGAGGACCCAGGGTTCCCGAGGGAACACCTCTGGAGGAGGAAC
GTCCAGCACCAAGCTGGAGACAGGCAGTGTCCCAATCCGGGTGTGGAACCAGAACTTCGCCTGGAGAACAGAG
AGCAACAAGTGTAAGGCTCCAAAGCCTGAAGGCTTCTCCCGCCACCCTTCCACCCCGGGTGGTGATCATGGC
AGAGGTGGCCCTGGTGGCATGTGGGGAGGAAGAGGTGGCTCATGGATCATGGTGGTCCCGGTGGAATGTTTACA
GGTGGCTGTGGTAGAGACAGAAGTGGCTTCTGTGGTGGCTGGGCATGGACCGAGGTGGCTTTGGTGGAGGAAGAC
AAGGTGGCCCTGGGGGGCCCCCGGACCTTTGATGTAACCAATGGGAGGAAGAAGAGGAGGACGTGAAGGACCTGG
AAAAACGGATAAAGCGAGCACTGTTCAGGAGCGCAGAGATCAGCCCTACTAGATGCAGAGAACCCGCAGAGCTGC
ATTGACTACCAGATTTATTTTTTAAACCAGAAAATGTTTTAAATTTATAATTCCATATTTATAATGTTGGCCACA
ACATTATGATTATTTCTTCTCTGTACTTTAGTATTTTTACCATTTGTGAAGAAACATTAAAAACAAGTTAAATGG
TA

527/6881
FIGURE 491

MGVYGQESGGFSGLGENRSMSPDNWGRGRGGFDRGGMSRGGGGCGGMSAGEQVGFNKPGGPMDEGPDLDLG
PPVDPDESDNSAIYVQGLNDNVTLDDLVDFFKQCGVVKMNRTEQPMIHTYLDKETRKPKGDATVSCEDSPTAK
AAVEWFDGKDFQGSKLKVS LARKRPPVNSMQGGMPPHEGRGMPPPLCGGPGGPGSPGGPMGHMGGRGDRGGLPP
RGPQGSRGNTSGGGNVQHQAQDRQCPNPGCGNQNFARWTESNKCKAPKPEGFLPPFFPPPGGDHGRGGPGGMWGG
RGGLMDHGPGGMFRGGCGRDRSGFCGGWAWTEVALVEEDKVALGGPRTFDVTNGRKKRRT

528/6881

FIGURE 492

AACGAGGAGGAAGGAGAGAAAATGGCATCCATGGATTACAGTACCTATAGCCAAGCTGCAGCGCAGCAGGGCTAC
AGTGCTTACAACACCCAGCCCACTCAAGGATATGCACAGCGCCTATGCAACTTCTTATGGACAGCCTCCCACTGG
TTATACTACTCCAAGTGGCCCCCAGGCATACAGCCAGCCTGTCCAGGGGTATGGCACTGGTGCTTATGATACCAC
CACTGCTACAGTCACCACCACCCAGGCCTCCTATGCAGCTCAATCTGCATATGGCACTCAGCCTGCTTATCCAGC
TTATGGGCAGCAGCCAGTAGCCACTGCACCTACAAGACTGCAGAATGGAAACAAGCCCACTGAGACTAGTCAACC
TCAATCTAGCACAGGGGGTTACAACCAGCCAGCCTAGGATATGGACAGAGTAACTGCAGTTATCCCCAGGTACC
TGGGAGCTACTCCATGCAGCCAGTCACCGCACCTCCATCCTACCCTCCTACCAGCTATTCTCTACACAGCCAAC
TAGTTATGATCAGAGCAGTTACTCTCAGCAGAACACCTATGGGAAACCGAGCAGCTATGGACAGCAGAGTAGCTA
TGGTCAACAAAGCAGCTATGGGCAGCTGCCTCCCACTAGTTACCCACCCCAAAGTTGATCCTACAGCCAAGCTCC
AAGTCAGTATAGCTAACAGAGCAGCAGCTACGGGCAGCAGAGTTCATTCTGACAGGACCACCCAGTAGCATGGG
TGTTTATGGGCAGGAGTCTGGAGGATTTCCGGACTAGGAGAGAACCGGAGCATGAGTGGCCCTGATAACTGGGG
CAGGGGAAGAGGGGGATTG

529/6881
FIGURE 493

AACGAGGAGGAAGGAGAGAAAATGGCATCCATGGATTACAGTACCTATAGCCAAGCTGCAGCGCAGCAGGGCTAC
AGTGCTTACAACACCCAGCCCCTCAAGGATATGCACAGACCACCCAGGCATATGGGCAACAAAGCTATGGAACC
TATGGACAGCCCATTGATGTCAGCTATACCCAGGCTCAGACCACTGCAATCTATGGGCAGACCGCCTATGCAACT
TCTTATGGACAGCCTCCCACTGGTTATACTACTCCAAGTCCCCCAGGCATACAGCCAGCCTGTCCAGGGGTAT
GGCACTGGTGCTTATGATACCACCACTGCTACAGTCACCACCACCCAGGCCTCCTATGCAGCTCAATCTGCATAT
GGCACTCAGCTGCTTATCCAGCTTATGGGCAGCAGCCAGTAGCCACTGCACCTACAAGACTGCAGAATGGAAAC
AAGCCCACTGAGACTAGTCAACCTCAATCTAGCACAGGGGGTTACAACCAGCACCAGCTGGAGACAGGCAGTGT
CCCAATCCGGGTGTGGAAACCAGAACTTCGCCTGGAGAACAGAGAGCAACAAGTGTAAGGCTCCAAAGCCTGAA
GGCTTCCTCCCGCCACCCTTCCCACCCCGGGTGGTGATCATGGCAGAGGTGGCCCTGGTGGCATGTGGGGAGGA
AGAGGTGGC

530/6881
FIGURE 494

GGAGGAAGGAGAGAAAATGGCATCCATGGATTACAGTACCTATAGCCAAGCTGCAGCGCAGCAGGGCTACAGTGC
TTACAACACCCAGCCCCTCAAGGATATGCACAGACCACCCAGGCATATGGGCAACAAAGCTATGGAACCTATGG
ACAGCCCATTGATGTCTAGCTATACCCAGGCTCAGACCACTGCAATCTATGGGCAGACCGCCTATGCAACTTCTTA
TGGACAGCCTCCCACTGGTTATACTACTCCAAGTGCAGGCTACAGCCAGCCTGTCCAGGGGTATGGCAC
TGGTGCTTATGATACCACTGCTACAGTCACCACCACCCAGGCCTCCTATGCAGCTCAATCTGCATATGGCAC
TCAGCCTGCTTATCCAGCTTATGGGCAGCAGCCAGTAGCCACTGCACCTACAAGACTGCAGAATGGAACAAGCC
CACTGAGACTAGTCAACCTCAATCTAGCACAGGGGGTTACAACAGCCAGCCTAGGATATGGACAGAGTAACTG
CAGTTATCCCCAGGTACCTGGGAGCTACTCCATGCAGCCAGTCACCGCACCTCCATCCTACCTCCTACCAGCTA
TTCCTCTACACAGCCAACTAGTTATGATCAGAGCAGTTACTCTCAGCAGAACACCTATGGGCAGCTGCCTCCAC
TAGTTACCCACCCAACTTGATCCTACAGCCAAGCTCCAAGTCAGTATAGCTAACAGAGCAGCAGCTACGGGCA
GCAGAGTTTATTCTGACAGGACCACCCAGTAGCATGGGTGTTTATGGGCAGGAGTCTGGAGGATTTCCGGACT
AGGAGAGAACCGGAGCATGAGTGGCCCTGAT

531/6881

FIGURE 495

TTTCCGGTGCTATGCCCCGGAAGCGGAAGTGCATCTTCGGGCTGTCAGAGTTGGTCTGTTACTCGGTGGTGGCG
GAGTCTACGGAAGCCGTTTTTCGCTTCACTTTTCCTGGCTGTAGAGCGCTTTCCCCCTGGCGGGTGAGAGTGCAGA
GACGAAGGTGCGAGATGAGCACTATGTTTCGCGGACACTCTCCTCATCGTTTTTATCTCTGTGTGCACGGCTCTGC
TCGCAGAGGGCATAACCTGGGTCCTGGTTTACAGGACAGACAAGTACAAGAGACTGAAGGCAGAAGTGAAAAAC
AGAGTAAAAAATTGGAAAAGAAGAAGGAAACAATAACAGAGTCAGCTGGTCGTCTATCAATGGTTCGAATGAAAT
CCATGTTTGCTATTGGCTTTTGTCTTACTGCCCTAATGGGAATGTTCAATTCCATATTTGATGGTAGAGTGGTGG
CAAAGCTTCCTTTTACCCCTCTTCTTACATCCAAGGACTGTCTCATCGAAATCTGCTGGGAGATGACACCACAG
ACTGTTCCCTTCATTTTCCTGTATATTCTCTGTACTATGTTCGATTTCGACAGAACATTCAGAAGATTCTCGGCCTTG
CCCCCTTCACGAGCCGCCACCAAGCAGGCAGGTGGATTTCTTGGCCCACCACCTCCTTCTGGGAAGTTCTCTTGAA
CTCAAGAACTCTTTATTTTCTATCATTCTTTCTAGACACACACATCAGACTGGCAACTGTTTTGTAGCAAGAG
CCATAGGTAGCCTTACTACTTGGGCCTCTTTCTAGTTTTGAATTATTTCTAAGCCTTTTGGGTATGATTAGAGTG
AAAATGGCAGCCAGCAAACCTTGATAGTGCTTTTGGTCCTAGATGATTTTTATCAAATAAGTGGATTGATTAGTTA
AGTTCAGGTAATGTTTATGTAATGAAAAACAAATAGCATCCTTCTTGTTTCATTTACATAAGTATTTTCTGTGGG
ACCGACTCTCAAGGCACTGTGTATGCCCTGCAAGTTGGCTGTCTATGAGCATTTAGAGATTTAGAAGAAAAATTT
ACAAGCCAAATACATGACATAAGATCAATAAAGAGGCCAAATTTTTAGCTGTTTTATGTACAAGGAGAGATCTGT
TTCATTTTGTGTTTGGCGTATTTCTAGATATAAGTTTTAGCATGGGCCAGGAAGGACTAAAAATAAAAGTTTTTAAG
GTA CTG

532/6881
FIGURE 496

MSTMFADTLLIVFISVCTALLAEGITWVLVYRTDKYKRLKAEVEKQSKKLEKKKETITESAGRQKKKKIERQEEK
LKNNNRDLMSVRMKSMFAIGFCFTALMGMFNSIFDGRVVAKLPFTPLSYIQGLSHRNLLGDDTTDCSFIFLYILC
TMSIRQNIQKILGLAPSRAATKQAGGFLGPPPPSGKFS

533/6881
FIGURE 497

AGACAGTTTTGAAGTTTTCAAAGACTGGCTCTGCTGTTAAGAAGTTGTACTTAAAGCGGAGGAGCTAAGCCACCT
GCCAAAATGTGCAAAGGACTTGCAGCTTTGCCCCACTCATGCCTGGAAAGGGCCAAGGAGATTAAGATCAAGTTG
GGAATTCCTCTCCAGAAGCCAGACTCAGTTGGTGACCTTGTCTATTCCGTACAATGAGAAGCCAGAGAAACCAGCC
AAGACCCAGAAAACCTCGCTGGACGAGGCCCTGCAGTGGCGTGATTCCCTGGACAAACTCCTGCAGAACAACTAT
GGACTTGCCAGTTTCAAAGTTTTCTGAAGTCTGAATTCAGTGAGGAAAACCTTGAGTTCTGGATTGCCTGTGAG
GATTACAAGAAGATCAAGTCCCTGCCAAGATGGCTGAGAAGGCAAAGCAAATTTATGAAGAATTCATTCAAACG
GAGGCTCCTAAAGAGGTGAATATTGACCACTTCACTAAGGACATCACAATGAAGAACCTGGTGGAACCTTCCCTG
AGCAGCTTTGACATGGCCCAGAAAAGAATCCATGCCTGATGGAAGGATTCTCTGCCTCGCTTTGTGCGCTCT
GAGTTTTATCAGGAGTTAATCAAGTAGTAATTTAGCCAGGCTATGAAATCATCCTGTGAGTTATTTCCCTCCATAA
TAACCTGCAATTTCCCATTAATCTACATATCTTCCACAGCAGCTTTGCTCAGTGATACCCACATGGGAAAAATC
CCAGGGGATGTGCTTACTCTTTTTGCCACACIGCTTTGGATACTTATCTACTGTCCGAAGGCCTTCTTTCCCC
ACTCAATTCCTCTGCCCTGTTATTAATTAAGATATCTTCAGCTTGTAGTCAGACCCAATCAGAATCACAGAAAA
ATCCTGCCTAAGGCAAAGAAATATAAGACAAGACTATGATATCAATGAATGTGGGTTAAGTAATAGATTTCCAGC
TAAATTGGTCTAAAAAGAATATTAAGTGTGGACAGACCTATTTCAAAGGAGCTTAATTGATCTCACTTGTTTTA
GTTCTGATCCAGGAGATCACCCCTCTAATTATTTCTGAACCTTGGTTAATAAAAGTTTTATAAGATTTTTATGAAG
CAGCCACTGTATGATATTTAAGCAAATATGTTATTTAAATATTTGATCCTTCCCTTGGACCACCTTCATGTTAG
TTGGGTATTATAAATAAGAGATACAACCATGAATATATTATGTTTATACAAAATCAATCTGAACACAATTCATAA
AGATTTCTCTTTTATACCTTCCTCACTGGCCCCCTCCACCTGCCCATAGTCACCAAATCTGTTTTAAATCAATG
ACCTAAGATCAACAATGAAGTATTTTATAAATGTATTTATGCTGCTAGACTGTGGGTCAAATGTTTCCATTTTCA
AATTATTTAGAATTCTTATGAGTTTAAAATTTGTAAATTTCTAAATCCAATCATGTAAATGAAACTGTTGCTCC
ATTGGAGTAGTCTCCACCTAAATATCAAGATGGCTATATGCTAAAAAGAGAAAATATGGTCAAGTCTAAATGG
CTAATTGTCCTATGATGCTATTATCATAGACTAATGACATTTATCTTCAAACACCAAATTTGCTTTAGAAAAAT
TAATGTGATTACAGGTAGAGGCCTTCTAGGTGAGACACTTTAAGGTACACTGCATTTTGCAGAAAAAAAAAAAA
AAAAGTAATCTTTTAGCAACCCAGTATTCCTTCACTATTTGCTTCCCTGCATTAGCAAATTTTACTTACAGTCA
AAAGTGCAGATTTTACTCCTGACGTGTCTCATTACAGCTAAATAATAGGCCATAGGACTTTTGGTAGGTTTAA
ACTTTTAATTCGTATTTTATGATTATAAGTCTTGCTAGAATTTTTCTAATCTTTAGTAGATTGATTAAATAA
TGATTCACAGAATTTAGTAACAGAATCAAATAAGCCATGTATGAGGGTAATCGAGATGAGGATATTAACATAA
AGAAATAGGGTGATTTTTAAAGGATTAATAAAATTTCTGAAATGTTAAGTAG

534/6881

FIGURE 498

MCKGLAALPHSCLERAKEIKIKLGILLQKPDSVGDLVIPYNEKPEKPAKTQKTSIDEALQWRDSLDKLLQNNYGL
ASFKSFLKSEFSEENLEFWIACEDYKKIKSPAKMAEKAKQIYEEFIQTEAPKEVNIDHFTKDITMKNLVEPSLSS
FDMAQKRIHALMEKDSLPRFVRSEFYQELIK

535/6881
FIGURE 499

GACACCTTTTAAAATGCAGAACTAACTGAGGCATTTTCAGTAACTTTGCTTTCAAATCAATAAAGTCAAATGTATG
GAAACATTTTGTGCCCTACTCTCCATACCCTGTGTACTCAAATTCTCTACTGTATGAATTATGCTTTAAGTAGAA
TTCAGTGCCAAGGAGAACTTGGTGAAATAAATTATTTTAAATTTTTTTTATCCTTTACAAAGCCATGGATTTTA
TTTGGTTGATGTGTGCTCTGTACACAAGCCATTTCAATAGGATGGAGCTGTTAATTATTTTCCAAAGAGTAATAG
ACATGCAAAAGTTTCAATTAATAAACTGGGCCATTAAACAAATAAATTAATAAACTAATAAGCATTCCCTTCTAGGTT
TTTGCCAACTGCCTATCCAATAACAAATTTGAGAATCGTTGAAAAAGCTAGTTATATTTTCAGAGAAATGATTTT
CATTATTGAACTGTTCTCCCTAGCAGGCCATTTTCCCTTTTTCTGGGAGTTTAGCAAGTTTAGGAGAGAATAG
TCATGAAAAGAAAGGGAAGAAAGGGGAGAAGGGAAGAGGTTAAAAAGTAAGTGCTCAGACCTATGAACGTAATCC
CTTTGCTAGAAATATTTAAGAGCAGCTCAGCTTGGTTGAAACTGAGTTTTGTCTATCTCCATATTTGCAGGAAGG
TATTTTCTGACTTGCAATGCAGCTAGATGTAAATTTTATTTTATCATCCTAGAAAGCCTTGACTAGAAAAATGA
ATAAATATTGAGGGTTTCTGTCCATATCTGGCTTGCAATGTGCCAGAAAGCAGAGAAATAGAAAATGTAATCTCCA
ACATCCAAGCATCGAAACCCAAGGGTAGGCAATTTCTATGTAGGTTTTGGACATGAAGTTGGTGATCTTGGTT
TATGCTGGCTCAACTGCTATTAAACCTCTCTGGCTTATAGTCTCTTCATTCTATTAGACAAGCACGTATCGAACA
CTTGCTTCGCACAAGGCTCTTTAGTTAACAATTTAGCAGCTACTGTTTGTGTTAAACACACTTTTCACCAAAATAG
GTTCTGAGGCAAACGAGAGCAATGACTATTTAAAGAAAGGCTTTCCAGCATCACTTACACATCCCAAACTAAA
AAGATCAACTCTTCCAAGTGAAGAAAGACTCCTGGCTTTGAATGGAACTTACAGCAGAGAGTCACAGGCCACGG
CAACAACAACGACAACAACAACAAATTTGGAATATTATTCTCAACTCACGTTTTAATAATACATCTTAATTATTTT
TCTAGTAGAGAACTACAAATCAGCCTCTTCAACATTTATATACAGTTTAATAAGCCTCTTGCAAGTTACTTGTT
CTCTCACCTGAGGTATTTTTTCTCCCACTTGCCCCCTGTTCCCTCCCTTCTCTCCCTTGCAAGAGGAA
ATATTTAACATATTTGGGTCCAATTCATAATGTAATAATTAATACATTTAAAGCATTTAACCTCCTTTCTAGA
AAAATGCACAGGCTAAGGCATAGACAAAACAAAGAGAAATGCTGAGAAATTTGCCACTGGAGACAAGCAATCTGA
ATAAATATTTGCCAAAAGTTCTTTTATGTATATAGTGTGAGGATTTGAAGGAGCTATTTTTTTTAAATGTTGCA
ACTAGCAACTCATCTTCGGAAGACACAGCCAGGAGAATGAAGTAGAAGTGAAAGGTTTATAAATCCATTTGTAAG
CATTATCCCATATATTTTAAATTCAAGAAAAATTGTGTTTATCTTTAGAATTTGTATTCAATACCTTTATGTAC
TATGTGACTCATGCTTCTGGATAAATAAAGCACCAAAATATGTATCTGTAACCACAATCACACATATTATATTA
TATATATCTATATAACAAAAAAAAAAAAAAAAAAAA

536/6881

FIGURE 500

MYGNILCPTLHTLCTQILYCMNYALSRIQCQGEINYNFFFIYKAMDFIWLMCALYTSHFNRMELLIIFQR
VIDMQKFQ

537/6881
FIGURE 501

AGCGGCCCCCTGCAGAGGCAGGTCCACCTGAGAGGCCGGCCGGCCTCCCAGCCCAGTGTTCATCCGGGGCATCACCT
ACTATAAAGCCAAGGTCTCTGAAGAAGAGAATGACATTGAAGAGCAGCAAGATGAGTTTTTCAGCGGTGACAATG
GAGTGGATTGTGCTGATTGAAGATCAGCTCCTGAGACACAACGGCCTGATGACCAGTGTACCCGGAGGCCATGCAG
CCACCCGTCAGGGACACAGCACTGCTGTGACAAGCGACCTGAACGCTCGGACCGCACCTGGTCTCTCAGCACTGC
CACAGCCCTCGACCTCAGATCCCAGCATCGCCAACCATGCCTCAGTGGGACCAACACTCCAAACAACCTCGGTGT
CTCCAGATCCCACAAGGGAGTCAGTCCTGCAGCCTTCTCCTCAGGTACCAGCCACCACTGTGGCCACACAGCCA
CCCAGCAACCAGCAGCCCCAGCTCCTCCGGCAGTGTCTCCAGGGAGGCATTGATGGAAGCTATGCACACAGTCC
CAGTGCCTCCCACCACAGTCAGAACAGACTCGCTGGGGAAGATGCTCCTGCTGGGTGGGGAACAACCCCTGCCA
GCCCCACGCTGAGCCCCGAAGAAGATGACATCCGGAATGTCTATAGGAAGGTGCAAGGACACTCTCTCCACAA
TCACGGGGCCGACCACCCAGAACACATATGGGCGGAATGAAGGGGCCTGGATGAAGGACCCCTGGCCAAGGATG
AGCGGATTTTACGTAACCAACTATTACTACGGCAACACCCCTGGTAGAGTTCCGGAACCTGGAGAAGTTCAAACAAG
GTCGCTGGAGCAATTCTACAAGTCCCGTACAGCTGGATCGGCACAGGCCACGTGGTATACAATGGCGCCTTCT
ACTACAATCGCGCCTTACCCGCAACATCATCAAGTACGACCTGAAGCAGCGCTACGTGGCTGCCTGGGCCATGC
TGCAATGACGTGGCCTACGAGGAGGCCACCCCTGGCGATGGCAGGGCCACTCAGACGTGGACTTTGCTGTGGACG
AGAATGGCCTATGGCTCATCTACCCGGCCCTGGACGATGAGGGCTTCAGCCAGGAGGTCAATGTCCTGAGCAAGC
TCAATGCCGCGGACCTGAGCACACAGAAGGAGACCACATGGCGCACGGGGCTCCGGAGGAATTTCTACGGCAACT
GCTTCGTATCTGTGGGGTGTGTATGCCGTGGATAGCTACAACCAGCGGAATGCCAACATCTCCTACGCTTTCG
ACACCCACACCAACACACAGATCGTCCCCAGGCTGCTGTTTCGAGAATGAGTATTCCTATACGACCCAGATAGACT
ACAACCCCAAGGACCGCCTGCTCTATGCCTGGGACAATGGCCACCAGGTCACTTACCATGTCATCTTTGCCTACT
GACACCCCTTGTCACCCACAAGCAGAGGACAGAGGGGTCACTAGCACCTTGTTGTGTATGTGTGTGCGCGCACGTGT
GTGTAGGTGGGTATGTGTTGTTTTAAAAATATATATTTTGTATAATATTGCAAATGTAAAAATGACAATTTGGG
TCTATTTTTTTTATATGGATTGTAGATCAATCCATACGTGTATGTGTGCTGGTCTCATCCTCCCCAGTTTATATTTT
GTGCAAATGAACTTCTCCTTTTGACCAGTAACCACCTTCCTTCAAGCCTTCAGCCCCCTCCAGCTCCAAGTCTCAG
ATCTCGACCATGAAAAGGTTTCTTCATCTGGGTCTTGCAGGAGGCAGGCAACACCAGGAGCAGAAATGAAAGAG
GCAAGAAAGAAGTGCTATGTGGCGAGAAAAAAGTTTAAATGTATTGGAGAAGTTTTAAAAAACCCAGAAAAACG
CTTTTTTTTTTTAATAAAGAAGAAATTTAAATC

538/6881
FIGURE 502

MTSVTRRPAATRQGHSTAVTSDLNARTAPWSSALPQPSTSDPSIANHASVGPTLQTTSVSPDPTRESVLQPSPQV
PATTVAHTATQQPAAPAPPAVSPREALMEAMHTVPVPPTTVRTDSLKGDAPAGWGTTTPASPTLSPEEEDDIRNVI
GRCKDTLSTITGPTTQNTYGRNEGAWMKDPLAKDERIYVTNYYYGNTLVEFRNLENFKQGRWSNSYKLPYSWIGT
GHVVYNGAFYYNRAFTRNIIKYDLKQRYVAAWAMLHDVAYEEATPWRWQGHSDVDFAVDENGLWLIYPALDDEGF
SQEVIVLSKLNAADLSTQKETTWRITGLRRNFYGNCFVICGVLYAVDSYNQRNANISYAFDTHNTQIVPRLLFEN
EYSYTTQIDYNPKDRLLYAWDNGHQVTYHVIFAY

539/6881
FIGURE 503

GGCACGAGGGATGCAAGGAGATGAGACAGTTAGATTTACTTCCTCTTTTCTAATCTGAGAGGTTTCATGTTGAAG
AAAATCAGTGTTGGGGTTGCAGGAGACCTAAACACAGTCACCAATGAAGCTGGGCTGTGTCCTCATGGCCTGGGCC
CTCTACCTTTCCCTTGGTGTGCTCTGGGTGGCCAGATGCTACTGGCTGCCAGTTTGGAGACGCTGCAGTGTGAG
GGACCTGTCTGCACTGAGGAGAGCAGCTGCCACACGGAGGATGACTTGACTGATGCAAGGGAAGCTGGCTTCCAG
GTCAAGGCCTACACTTTCAGTGAACCTTCCACCTGATTGTGTCTATGACTGGCTGATCCTCCAAGGTCCAGCC
AAGCCAGTTTTTGAAGGGGACCTGTCTGGTTCTGCGCTGCCAGGCCTGGCAAGACTGGCCACTGACTCAGGTGACC
TTCTACCGAGATGGCTCAGCTCTGGGTCCCCCGGGCCTAACAGGGAATTCTCCATCACCGTGGTACAAAAGGCA
GACAGCGGGCACTACCACTGCAGTGGCATCTTCCAGAGCCCTGGTCTGGGATCCAGAAACAGCATCTGTTGTG
GCTATCACAGTCCAAGAAGTGTTCAGCGCCAATTCTCAGAGCTGTACCCTCAGCTGAACCCCAAGCAGGAAGC
CCCATGACCCTGAGTTGTGAGACAAAGTTGCCCTGCAGAGGTGAGCTGCCCGCCTCCTCTTCTCCTTCTACAAG
GATGGAAGGATAGTGCAAAGCAGGGGGCTCTCCTCAGAATTCCAGATCCCCACAGCTTCAGAAGTCACTCCGGG
TCATACTGGTGTGAGGCAGCCACTGAGGACAACCAAGTTTGGAACAGAGCCCCAGCTAGAGATCAGAGTGCAG
GGTGCTTCCAGCTCTGCTGCACCTCCCACATTGAATCCAGCTCCTCAGAAATCAGCTGCTCCAGGAAGTCTCCT
GAGGAGGCCCCCTGGGCCTCTGCCTCCGCCGCCAACCCCATCTTCTGAGGATCCAGGCTTTTCTTCTCCTCTGGGG
ATGCCAGATCCTCATCTGTATCACCAGATGGGCCTTCTTCTCAAACACATGCAGGATGTGAGAGTCTCCTCGGT
CACCTGCTCATGGAGTTGAGGGAATTATCTGGCCACCAGAAGCCTGGGACCACAAAGGCTACTGCTGAATAGAAG
TAAACAGTTCATCCATGATCTCACTTAACCACCCCAATAAATCTGATTCTTTATTTCTCTTCTCTGCTGCACA
TATGCATAAGTACTTTTACAAGTTGTCCCAGTGTTTTGTTAGAATAATGTAGTTAGGTGAGTGTAATAAAATTTA
TATAAAGTGAGAATTAGAGTTTAGCTATAATTGTGTATTCTCTCTTAACACAACAGAATTCTGCTGTCTAGATCA
GGAATTTCTATCTGTTATATCGACCAGAATGTTGTGATTTAAAGAGAACTAATGGAAGTGGATTGAATACAGCAG
TCTCAACTGGGGGCAATTTTGCCCCCAGAGGACATTGGGCAATGTTTGAGACATTTTGGTCATTATACCTGGG
GGGTGGGGGATGGTGGGATGTGTGTGCTACTGGCATCCAGTAAATAGAAGCCAGGGGTGCCGCTAAACATCCTA
TAATGCACAGGGCAGTACCCACAAACGAAAAATAATCTGGCCCAAAATGTCAGTTGTACTGAGTTTGAGAAACCC
CAGCCTAATGAAACCCTAGGTGTTGGGCTCTGGAATGGGACTTTGTCCCTTCTAATTATTATCTCTTTCCAGCCT
CATTCAGCTATTCTTACTGACATACCAGTCTTAGCTGGTGCTATGGTCTGTTCTTTAGTTCTAGTTTGTATCCC
CTCAAAAGCCATTATGTTGAAATCCTAATCCCCAAGGTGATGGCATTAGAAGTGGGCCTTTGGGAAGTGATTAG
ATCAGGAGTGCAGAGCCCTCATGATTAGGATTAGTGCCTTATTTAAAAAGGCCCCAGAGAGCTAACTCACCCCTT
CCACCATATGAGGACGTGGCAAGAAGATGACATGTATGAGAACCAAAAAACAGCTGTGCGCAAACACCGACTCTG
TCGTTGCCTTGATCTTGAAGTTCAGCCTCCAGAAGTATGAGAAATAAAATCTGTTGTTTGTAAAAA
AAAAAA

540/6881
FIGURE 504

MKLGCVLMAWALYLSLGLWVAQMLLAASFETLQCEGPVCTEESSCHTEDDLTDAREAGFQVKAYTFSEPFHLIV
SYDWLILQGPAKPVFEGDLLVLRCAWQDWPLTQVTFYRDGSALGPPGPNREFSITVVQKADSGHYHCSGIFQSP
GPGIPETASVVAITVQELFPAPILRAVPSAEPQAGSPMTLSCQTKLPLQRSAARLLFSFYKDGRIVQSRGLSSEF
QIPTASEDHSGSYWCEAATEDNQVWKQSPQLEIRVQGASSSAAPPTLNPAPQKSAAPGTAPEEAPGPLPPPPTPS
SEDPGFSSPLGMPDPHLYHQMGLLLKHMQDVRVLLGHLLMELRELSGHQKPGTTKATAE

541/6881
FIGURE 505

AAGCAGTGGTATCAACGCAGAGTGCCCATACGGCCGGGGTGGCATCATGTGGCAGCTGCTCCTCCCAACTGCT
CTGCTACTTCTAGTTTCAGCTGGCATGCGGACTGAAGATCTCCCAAAGGCTGTGGTGTTCCTGGAGCCTCAATGG
TACAGGGTGCTCGAGAAGGACAGTGTGACTCTGAAGTGCCAGGGAGCCTACTCCCCTGAGGACAATTCACACAG
TGGTTTCACAATGAGAGCCTCATCTCAAGCCAGGCCTCGAGCTACTTCATTGACGCTGCCACAGTCGACGACAGT
GGAGAGTACAGGTGCCAGACAAACCTCTCCACCCTCAGTGACCCGGTGCAGCTAGAAGTCCATATCGGCTGGCTG
TTGCTCCAGGCCCTCGGTGGGTGTTCAAGGAGGAAGACCCTATTACCTGAGGTGTCACAGCTGGAAGAACACT
GCTCTGCATAAGGTACATATTTACAGAATGGCAAAGGCAGGAAGTATTTTCATCATAATTCTGACTTCTACATT
CCAAAAGCCACACTCAAAGACAGCGGCTCCTACTTCTGCAGGGGGCTTGTTGGGAGTAAAAATGTGTCTTCAGAG
ACTGTGAACATCACCATCACTCAAGGTTTGGCAGTGTCAACCATCTCATCATTCTTTCCACCTGGGTACCAAGTC
TCTTTCTGCTTGGTGATGGTACTCCTTTTTTGCACTGGACACAGGACTATATTTCTCTGTGAAGACAAACATTCTGA
AGCTCAACAAGAGACTGGAAGGACCATAAATTTAAATGGAGAAAGGACCCTCAAGACAAATGACCCCCATCCCAT
GGGGGTAATAAGAGCAGTAGCAGCAGCATCTCTGAACATTTCTCTGGATTTGCAACCCTATCATCCTCAGGCCTC
TCTACAAGCAGCAGGAAACATAGAACTCAGAGCCAGATCCCTTATCCAACTCTCGACTTTTCCTTGGTCTCCAGT
GGAAGGGAAAAGCCCATGATCTTCAAGCAGGGAAGCCCCAGTGAGTAGCTGCATTCTAGAAATTGAAGTTTCAG
AGCTACACAAACACTTTTTCTGTCCCAACCGTTCCCTCACAGCAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA

542/6881
FIGURE 506

MWQLLLPTALLLLVSAGMRTE~~L~~PKAVVFLEPQWYRVLEKDSVTLKCQGAYSPEDNSTQWFHNESLISSQASSYF
IDAATVDDSGEYRCQTNLSTLS~~D~~PVQLEVHIGWLLQAPRWVFKEEDPIHLRCHSWKNTALHKVTYLQNGKGRKY
FHHNSDFYIPKATLKDSGSYFCRGLVGSKNV~~S~~SETVNITITQGLAVSTISSFFPPGYQVSFCLVMVLLFAVDTGL
YFSVKTNIRSSTRDWKDHKFKWRKDPQDK

543/6881
FIGURE 507

ATGGCGGACGGGACCTGTCAGGACGTGGCCATCGTGGGCTACAAGGACTCGCCCTCCATCTGGGCCGCGTCCCC
GGGAAAACCTTCGCCAACATCGCGCCAGCTGAGGTTGGTGTCTGGCTGGCAAAGACAGGTCAAGATGGGGAATT
TTGCATGGATCTTCGTATACCAAGAGTACCGGTGGAGCCCCACCTTCAATGTCACTGTCACCAAGACTGACAAG
ACGCTAGTCCTGCTGATGGGCAAAGAAGGTGTCCACGGTGGTTTGATCAACAAGAAATGTTATGAAATGGCCTCC
CACCTTCGGCGTTCCAGTACTGA

544/6881
FIGURE 508

MADGTCQDVAIVGYKDSPSIWAAVPGKTFANIAPAEVGVLAGKDRSRWGILHGSSYTKSTGGAPT FNVTVTKTDK
TLVLLMGKEGVHGGLINKKCYEMASHLRRSQY

545/6881
FIGURE 509

ATGTTGTAGGATTCAGTGGTGAATCCTTCCACCCAGGCCTTGCTGCTGCAGCCTCTGCCAGAGCCAGCTCC
TGGTGGACCCATGTGGAAATGGGACCTCCAGATCCCATCCTGGGAGTCACTGAAGCCTTTAAGAGGGACATCAAT
AGCAAAAAGATGAATCTGGGAGTTGGGATGATAATGGGAAGCCGCTTTGCCAGTGGTGATGGTGATAAGGATGCC
TGGGCTCTGCACCACTTCATCGAACAGGGCATTAAATGTTTGTCTCTGCCAATCATATGCCAAGAACATGGGCTTA
TATGGTGAGCATGTAGGAGCCTTCACTGTGGTCTGCAAAGATGCAGATGAAGCCAAAAGGGTAGAGTCACAGTTG
AAGATCTTGATTAGTCCCATGTATTCCAACCTCTCCCTCAATGGGGCCCAGATTGCTTCTACCATTCCAAACACC
CCAGTTGGTTTGAAGGTCTGGGTCTGGGAGGCCTACATGATCATCCTCAGGTTATGAGTGACGAGAAAGAGACA
ACACTGAGAGCAGTGGCTGTCAAGTACTGGATCAGAGTCCGGCTGTGCCACTTCCTTGAAACATTATGTCTCCG
GACCTTACTGGCTGGAAAAAACTGCAGCAGGAAGATGGCCTTACCTCACTTCTAGCTTCCAAGTCTCACAAGAGA
GCACAGGCAGAACCGAAGCCACATCCAGGACCTGGGCTGAGAACATCTGGCACATGGAGCCTGTAG

546/6881
FIGURE 510

MVVGFTFESFHPGLAAAASARASSWTHVEMGPPDPILGVTEAFKRDINSKKMNLGVGMIMGSRFASGDGDKDA
WALHHFIEQGINVCLCQSYAKNMGLYGEHVGAFTVVCKDADEAKRVESQLKILISPMYSNSPLNGAQIASTIPNT
PVGLKVWVMGGLHDPQVMSDEKETTLRAVAVKYWIRVRLCHFLETFMSPDLTGWKKLQQEDGLTSLLASKSHKR
AQAEPKPHPGPGLRTSGTWSL

547/6881

FIGURE 511

CGGTTGCTTGCTGTCCCAGCGGCGCCCCCTCATCACCGTCGCCATGCCCGGAGGTCTGCTTCTCGGGGACGTGGC
TCCCAACTTTGAGGCCAATACCACCGTCGGCCGCATCCGTTTCCACGACTTTCTGGGAGACTCATGGGGCATTCT
CTTCTCCCACCTCGGGACTTTACCCAGTGTGCACCACAGAGCTTGGCAGAGCTGCAAAGCTGGCACCAGAATT
TGCCAAGAGGAATGTTAAGTTGATTGCCCTTTCAATAGACAGTGTGAGGACCATCTTGCTGGAGCAAGGATAT
CAATGCTTACAATTGTGAAGAGCCACAGAAAAGTTACCTTTTCCCATCATCGATGATAGGAATCGGGAGCTTGC
CATCCTGTTGGGCATGCTGGATCCAGCAGAGAAGGATGAAAAGGGCATGCCTGTGACAGCTCGTGTGGTGTGTTGT
TTTTGGTCCTGATAAGAAGCTGAAGCTGTCTATCCTCTACCCAGCTACCACTGGCAGGAACCTTGTATGAGATTCT
CAGGGTAGTCATCTCTCTCCAGCTGACAGCAGAAAAAGGGTTGCCACCCAGTTGATTGGAAGGATGGGGATAG
TGTGATGGTCCTTCCAACCATCCCTGAAGAAGAAGCCAAAAAAGCTTTTCCCGAAAGGAGTCTTCACCAAGAGCT
CCCATCTGGCAAGAAATACCTCCGCTACACACCCAGCCTTAAGTCTCTTGGAGAAGTTGGTGTGTGAGCCAGA
GGATGTGAGCTGCCAATTGTGTTTTCTGTCAGCAATTCCATAAACACATCCTGGTGTATCACAGCCAAGGTTTT
TAGGTTGCTATACCAATGGCTTATTAAATGAAATGGCACTAAAAGTTTCTTGAGATTCTTTATACTCTCTGCCT
TCAGCAATCAATTCCATTACATACATCAGCACTCTGCTGGTTCTGTTTGAAATATGTTCTGTATTTAAACTCAA
TCTTGTGGATCTCTGTCAGGGCTTGTGACCAATGAAGTCATATTGTTGATGGTTGACAAAGCTTGCTTCACTCC
ATCAGAGAATGACTATCAATTTTTTTTTAACTGTCCTATCACGTCCTCTCCTGTACCCATTTGAAGAGTGGCA
GAACTTGAAGTTCAACTTCCTCTGTAAATATCCAAGTATAAAGCCAGGAACCTCTAGAATAACCCAGATGCGCT
TTAATTTTTTTTTAATATGTTTTGATCACAGAACTTCTAGAATAACCCAGATGCTCTTTCATATTCTTTTAATACA
TCTTGATCACAGCTGGGGGAAAAAAGCTTTTTAATTCTGTACCTTCCTAGTAGATAAGTGAAGAGCAGGGAAAG
AGACCTTTAAATATTTTGCTATAAAAAAATTTGTGATAAGTTTCTATCAAAATGGGGAGATTGCAGAAAAGGCTT
CCCTTGGCTCCCAAGGAGGTGTAGCAGGTGTGAGCAATATTAGTGCCATGTGCCTTTCACACAGGGTTTGCATTT
ATCAGTCTGTTTTCCGATGATGTGTACATGAAAGAGTACACCATGTGAAGAGAAGAGAGAAATGATTGAAAATGTT
TTAGTATAGAATCTTCTTGCAGTGGGTGCTATTTTCTAGATTTTACTTTTGTAGGAACAAAATAAAATCCTTT
GTT

548/6881

FIGURE 512

MPGGLLLGDVAPNFEANTTVGRIRFHDFLGDSWGILFSHPRDFTPVCTTELGRAAKLAPEFAKRNVKLIALSIDS
VEDHLAWSKDINAYNCEPTEKLPFPFIIDDRNRELAILLGMLDPAEKDEKGMPTARVVFVFGPDKKLKSILYP
ATTGRNFDEILRVVISLQLTAEKRVATPVDWKDGDSVMVLPTIPEEEAKKLFPGVFTKELPSGKKYLRYTPQP

549/6881

FIGURE 513

AGAGACCGTCCGAGGTAATTGTCTGCCACGAGTGCACATTCTGAAAACAGGAGATTTTAGTTCCTAAAAATGGGA
AGAACCTACATTGTAGAAAGAGACTGTTGGCCAGTATCTTTCAAACATAAATCTCCAAGGAAAGGCTTTTGTCTCT
GGCCTTTTAAATAGGACAGTGTTCGTACACAAAAGGATTATGTGATTCTTGCCACTAGAACGCCACCCAAAGAGGAG
CAAAGTGAGAACCTCAAACATCCCAAAGCTAAGTTGGATAACTTGGATGAAGAATGGGCCACAGAACATGCCTGC
CAGGTATCCAGAATGCTACCAGGGGGACTTTTAGTTCTTGGAGTATTTATTATTACTACTTTAGAACTGGCAAAT
GATTTTCAAATGCCCTGCGTAGACTAATGTTTGTGTGGAAAAGTCTATAAATAGAAAAGAGATTGTGGAATTTT
ACAGAGGAGGAAGTCTCAGAACGAGTGACACTTTCACATTTGTGCTTCTACAAAAAATATTTTGTCCGAACCTTAT
GATATCCATGATCCAAAGAGTTCAGCAAGACCAGCAGATTGGAAGTATCAAAGTGGATTATCATCCTCATGGCTT
TCTTTAGAGTGTACAGTTCACATTAATATTCACATCCCACTTTCTGCTACTTCTGTCTCAGCTATACTCTGGAGAAA
AATACAAAGAAATGGACTTACACGCTGGGCCAAGGAAATAGAAAATGGTGTATTTTATTGATTAATGGACAAGTTAA
GATGAAGATTGTGACCTATTAGAAGGACAGAAAAATCTTCTAGAGGAAATACTCAAGCAACTAGTCATTCTTTT
GATGTCAGAGTGCTAACGCAGTTGCTCCTGAATTTCAGACCACAGATCCACAGCCACAGTCCAGATATGTAGCGGT
TCTGTAAACCTTAAGGGTGCTGTGAAATGCAGAGCTTATATCCACAGCAGTAAACCCAAAGTTAAAGATGCTGTG
CAGGCAGTAAAGAGGGATATATTGAACACAGTTGCTGATCGTTGTGAAATGCTATTTGAGGATCTGCTTTTGAAT
GAAATTCAGAAAAAAGATTCTGAAAAAGAGTTCCACGTCCTCCCTTATCGAGTCTTTGTTCCCTTCCTGGA
TCCACTGTAATGTTGTGTGATTATAAATTTGACGATGAGTCAGCTGAAGAAATCAGGGACCATTTTATGGAGATG
TTGGATCACACAATTCAAATAGAAGATTTGGAAATTCAGAGGAAACAAACACAGCTTGATGAGTCTTCTATG
AATAGTCAAGCTTCATTGGACAACACAGATGATGAACAACCAAAACAACCAATTAAACTACAATGTTATTGAAA
ATTAGCAAAACATAGGTGTGATTGCAGCATTTACAGTTGCAGTCCTTGCTGCGGTATCTCCTTTTATTACTTC
AGTGATTAGGGTGAGGCACAAAGAGTTTCTTGATCATCCAGAGAACATTGACAGACAATTATGAATAATAAGAT
GTTAACAATCCATCTGTATTTAAACACTAGCAGCCAGATCTGCTGCCATGATGCCTATTTGGTGTGTTTCTGAT
TAAAATGAAATCACAAGCTGCCTTGTTTAGCCTGCTTTACATTGTAGGTGGCCCGCATTTCAGAAATAACGTTA
TGCATCTAGATGGAAGCTGCATGTAAACAAATCATTATTATCTATTTTAAAGCTTCAAATGATGGGATATGAT
CATAGATTTTAGTCTTACTAATCTGAATCACATATTAATCAGGACATTAAAACTTTAACAGAGGCATGATGGCT
CACACGTATAATCCTAATGCTTTGAGAGGCTGAGGTAGGAGCATCACCTGGGGCTGGGAGGGAGTTGGAGACCAG
CCTGGATGACATAATGAGATTCTGTCTCTACTGGAIAAAAAAAAAAAAAA

550/6881
FIGURE 514

MLPGGLLVLGVFIITTTLELANDFQNALRRLMFAVEKSINRKRLWNFTTEEVSERVTLHICASTKKIFCRTYDIHD
PKSSARPADWKYQSGLSLSSWLSLECTVHINIHIPLSATSVSYLEKNTKNGLTRWAKEIENG VYLINGQVKDEDC
DLLEGQKKSSRGNTQATSHSFDVRVLTQLLNSDHRSTATVQICSGSVNLKGAVKCRA YIHSSKPKVKDAVQAVK
RDILNTVADRCEMLFEDLLLNEIPEKKDSEKEFHVLPYRVFVPLPGSTVMLCDYKFDDESAEEIRDHFMEMLDHT
IQIEDLEIAEETNTACMSSSMNSQASLDNTDDEQPKQPIKTTMLLKIQQNIGVIAAFTVAVLAAGISFHYFSD

551/6881
FIGURE 515A

CGAGACGGAAGCGGGCTGGGAGGCGTCGGCGGCGGCAGCGCACGTGGTGACGTGCGAGGGGGTGCGGCGCGAGCG
GTCGGCGGCGGCGGAGGCAGTGTCTCCCGGTCGCGCGTGGAGGTCGGTTCGCTCAGAGCTGCTGGGCGCAGTTTCT
CCGCCTGCTGCTTCGGCGCGGCTGTATCGGCGAGCGAGCGAGTTCCCGCGAGTTCGCGTGGCGCTCCCCCTTCC
TTTCAGTCTCCACGGAAGTGGCCCTCGTCTTCTACTTGACCGTCCCGTCTTCCGCCGCTTCTGGCGCTTTCC
GTTGGGCGGATTCCCGCCGCTTCTCTGCTTCCCATCGAAGCTCTAGAAATGAATGTTTCCATCTCTTCAGAG
ATGAACCAGATTATGATGCATCATTATCACAGAAGAAATTCGTGTCTATAGCTTTTAAGGACTTGATTACATCAT
TTTCAAGCCTGATAGTTTTGGAAATCACCATTAGAGCTTAAGACACACCTGCCTTCATTTCAACCACCTGTCTTCA
TACCCTGACGAAGTGCACCTTTTAACACTCCTTTGTCCTTGGATTACTTAAGAGTTCAGAAATACATTTGCCA
CCAACAGAGTAGCCAAATTTATAAGGAAAAATGATTCCCAATGGATATTTGATGTTTGAGGATGAAAAATTTATT
GAGTCTTCTGTGCCAAATTAATGCCCTGAGGAAAAGTGGCCAGTTCTGTGATGTTGACTTCAGGTCTGTGGC
CATGAAATGTTAGCACACAGAGCAGTGTCTAGCTTGTGTCAGTCCCTATTTATTTGAAATCTTTAATAGTGATAGT
GATCCTCATGGAATTTCTCACGTTAAATTTGATGATCTCAATCCAGAAGCTGTTGAAGTCTTGTGTAATTATGCC
TACACTGCTCAGTTGAAAGCAGATAAGGAATTGGTAAAAGATGTTTATTCTGCAGCAAAAAAGCTGAAGATGGAT
CGAGTAAAGCAGGTTTGTGGTGATTATTTACTGTCTAGAATGGATGTTACCAGCTGCATCTCTTACCGAAATTTT
GCAAGTTGTATGGGAGACTCCCGTTTGTGTAATAAGGTTGATGCTTATATTCAGGAGCATTGTGTACAAATTTCT
GAAGAGGAGGAGTTTCTTAAGCTTCCAAGGCTAAAGTTGGAGGTAATGCTTGAAGATAATGTTTGCTTGGCCAGC
AATGGCAAATTTATATACAAAGGTAATCAACTGGGTGCAGCGTAGCATCTGGGAGAATGGAGACAGTCTGGAAGAG
CTGATGGAAGAGGTTCAAACCTTGTAATACTCAGCTGATCACAAGCTGCTTGATGGGAACCTACTAGATGGACAG
GCTGAGGTGTTTGGCAGTGATGATGACCACATTCAGTTTGTGTCAGAAAAAGCCACCACGTGAGAATGGCCATAAG
CAGATAAGTAGCAGTTCAACTGGATGTCTCTTCTCCAAATGCTACAGTACAAAGCCCTAAGCATGAGTGGAAA
ATCGTTGCTTCAGAAAAGACTTCAAATAACACTTACTTGTGCTGGCTGTGCTGGATGGTATATTCTGTGTCTATT
TTCTTTCATGGGAGAAAACAGCCCACAGAGCTCACCAACAAGTACTCCAAAATAAGTAAGAGTTTAAGCTTTGAG
ATGCAACAAGATGAGCTAATCGAAAAGCCCATGTCTCCTATGCAGTACGCACGATCTGGTCTGGGAACAGCAGAG
ATGAATGGCAAATCATAGCTGCAGGTGGCTATAACAGAGAGGAATGTCTCGAACAGTCGAATGCTATAATCCA
CATACAGATCACTGGTCTTTCTTGTCTCCCATGAGAACCAAGAGCCCGATTTCAAATGGCTGTACTCATGGGC
CAGCTCTATGTTGGTAGGTGGATCAAATGGCCACTCAGATGACCTGAGTTGTGGAGAGATGTATGATTCAAACATA
GATGACTGGATTCTGTTCAGAATTGAGAACTAACCGTTGTAATGCAGGAGTGTGTGCTCTGAATGGAAGTTA
TACATCGTTGGTGGCTCTGATCCATATGGTCAAAAAGGACTGAAAAATTGTGATGTATTGATCCTGTACAAAG
TTGTGGACAAGCTGTGCCCCCTCTTAACATTTCGGAGACACCAGTCTGCAGTCTGTGAGCTTGGTGGTTATTTGTAC
ATAATCGGAGGTGCAGAATCTTGAATTTGTCTGAACACAGTAGAACGATACAAATCCTGAAAATAATACCTGGACT
TTAATTGCACCATGAATGTGGCTAGGCGAGGAGCTGGAGTGGCTGTTCTTAATGGAAGTCTGTTGTATGTGGT
GGCTTTGATGGTTCTCATGCCATCAGTTGTGTGGAATGTATGATCCAAC TAGAAATGAATGGAAGATGATGGGA
AATATGACTTCACCAAGGAGCAATGCTGGGATTGCACTGTAGGGAACACCATTTATGCAGTGGGAGGATTCGAT
GGCAATGAATTTCTGAATACGGTGGAGTCTATAACCTTGAGTCAAATGAATGGAGCCCCATACAAAGATTTTC
CAGTTTTAAACAAATTTAAGACCCTCTCAAATAACAGGCTTAGTGATGTAATTATGGTTAGTAGAGGTACACTTG
TGAATAAGAGGGTGGGTGGGTATAGATGTGCTAACAGCAACACAAAGCTTTTGCATATTGCATACTATTAAAC
ATGCTGTACATACTTTTTGGGTTTTATTTGGAAAGGAATGCAAAGATGAAGGTCTGTTTTGTGTACTTTTAAAGACT
TTGGTTATTTTACTTTTTGGAAAAGAATAAACCAAGAATTGATTGGGCACATCATTTCAAGAAGTCCCTCTCCT
CCACATTTGTTTTGCCAATTTGCACATTAATGACTCTTCCCTCAAATGTGTACTATGGGGTAAAAGGGGTAGGG
TTTAAAGATGTAGACAGTTGGGTTTTTTAAGGGCCCTTTTTCAATAACTGGAACTCTATAACAAAGGATACTT
ATTTAAATAGATGACATTGACTATTTTTGTTTTATTAAAGGAAGCTTACATGCCTACCAATATTTAATCTTTT
ATGATTGCCTTTTTATAACTTTTTATATTCTCAGCAGAGTGCTTTACCAATTGAAGTAAATGTGGCAGGCTGGA
GTTATTGAAGCAGAGTGGCAGTCTTCAGTTGCAGAGTAGGGTCTGTCTTTTAACTCTGAGTGCAAACTTCAG
AGTTCTTGCTTGGCTGCAGTTTTTTTTCTTCAAGAATGCAGTACTAACATTTATTTGAGTGGAGTTACTGAACA
GTAACATAGCTGTGATTTTTGGTATTTGAAACACTGGTTTTAAATATTTTACTTGTGAGGGTATGTTTTATAT
AGCAAGACATTATATAGCAGTAAAAAATGGTGTTTTATCTTCTATATAAATCCTGTTTTTATTATTAACAAACA
GTCCTAAATAGCAGCCCTCAATTGTGAAAAAATTTACTTTAACTACATTAGGTTGTGAATGCAGGTTTTATCAG
AACTATGTTTTTGTTCAGTTTATCTGTTTCATATGGATAAATATTGGTTGGGATGACTTGGTGTCTAATGTGTAGT

552/6881

FIGURE 515B

GCTACACACCTAACTTATGGGGCCAAAATAGCATGTCCTAATGCTTGCTGCTGATTAAACACATTAAAGGTACT
TTGCAGGAAATCCTTGCACCATGGGATTAATATCCAATTGCTGCTTGTACACTCATTCACTAAAGTTTTGA
GAAATTTTTTTTTTCCAGTAATGAGCTTAAGAAATTTGTGGAAAATAACTCACCTGGCATCTTACATCTGAAATAA
GGAATGATATAAGGTTTTTTTTTCTCACAGAAGATGAAGCACACAGGAACCTAATGGGCCAACTGGGATGAGGTG
ACTATTCTGAGATGACTATTCAGTGGCTAACTTGGGTTAGGAAGAAAATAATTAGGTATTTCTCCAAATGTTCA
CTGGTACTCTGCCACTTTATTTCTCTCATCTGTTACACAAAGAACCACCAGGAAAGCAAATCAGTTTGGTTGGTA
ACTCTGTAATTCCTAACTATCACTGGTTTGGTTCTGGACTAAAACCTACATTGACAGATTGAATTTGCCTAATATG
ATGACTGTTTTTAATATGGATCTGTATGTGTTCTATTTCAGCACAAAGGAAATAAAATTTTAGTTGAGGATTCAGC

553/6881
FIGURE 516

MIPNGYLMFEDENFIESSVAKLNALRKSGQFCDVRLQVCGHEMLAHRAVLACCSPYLFEIFNSDSDPHGISHVKE
DDLNPEAVEVLLNYAYTAQLKADKELVKDVYSAKKLKMDRVKQVCGDYELSRMDVTSCISYRNFASCMGDSRLL
NKVDAYIQEHLLQISEEEFLKLPRLKLEVMLEDNVCLPSNGKLYTKVINWVQRSIWENGDSLEELMEEVQTLYY
SADHKLLDGNLLDGQAEVFGSDDDHQFVQKKPPRENGHKQISSSSTGCLSSPNATVQSPKHEWKIVASEKTSNN
TYLCLAVLDGIFCVIFLHGRNSPQSSPTSTPKLSKSLSFEMQQDELIEKPMSPMQYARSGLGTAEMNGKLIAGG
YNREECLRTVECYNPHDHSFLAPMRTPRARFQMAVLMGQLYVVGGSNHGSDDLSCGEMYDSNIDDWIPVPELR
TNRCNAGVCALNGKLYIVGSDPYGQKGLKNCDFDPVTKLWTSCAPLNIRRHQSAVCELGGYLYIIGGAESWNC
LNTVERYNPENNTWILIAPMNVARRGAGVAVLNGKLFVCGGFDGSHAISCVEMYDPTRNEWKMMGNMTSPRSNAG
IATVGNTIYAVGGFDGNEFLNTVEVYNLESNEWSPTYTKIFQF

554/6881
FIGURE 517

CCTTTGCCGCTGGTCGGGATTGGGATGTCGAAGAACACAGTGTCTGCGGCCCGCTTCCGGAAGGTGGACGTGGAT
GAATATGACGAGAACAAGTTTCGTGGACGAAGAAGATGGGGGCGACGGCCAGGCCGGGCGGACGAGGGCGAGGTG
GACTCCTGCCCTGCGGCAAGGAAACATGACAGCTGCCCTACAGGCAGCTCTGAAGAACCCCCCTATCAACACCAAG
AGTCAGGCAGTGAAGGACCGGGCAGGCAGCATTGTCTTGAAGGTGCTCATCTCTTTTAAAGCTAATGATATAGAA
AAGGCAGTTCAATCTCTGGACAAGAAATGGTGTGGATCTCCTAATGAAGTATATTTATAAAGGATTTGAGAGCCCG
TCTGACAATAGCAGTGCTATGTTACTGCAATGGCATGAAAAGGCACCTTGCTGCTGGAGGAGTAGGGTCCATTGTT
CGTGCTCTTGACTGCAAGAAAACTGTGTAGTCTGGCAGGAAGTGGATTATCTGCCTCGGGAGTGGGAATTGCTGG
TACAAAGACCAAAACAACCAATGCCACCGCTGCCCTGTGGGTAGCATCTGTTTCTCTCAGCTTTGCCTTCTTGC
TTTTTCATATCTGTAAAGAAAAAATTACATATCAGTTGTCTTTAATGAAAATTGGGATAATATAGAAGAAATT
GTGTTAAAATAGAAGTGTTCATCCTTTCAAACCATTTTCAGTGATGTTTATACCAATCTGTATATAGTATAATT
TACATTCAAGTTTAAATTGTGCAACTTTTAAACCCCTGTTGGCTGGTTTTTGTCTGTTTTGTTTTGTATTATTTT
TAACTAATACTGAGAGATTTGGTCAGAATTTGAGGCCAGTTTCTAGCTCATTGCTAGTCAGGAAATGATATTTA
TAAAAAATATGAGAGACTGGCAGCTATTAACATTGCAAACTGGACCATAATTTCCCTTATTTAATAAGCAAAATA
TGTTTTTGGAAATAAGTGGTGGGTGAATACCACTGCCAAGTTATAGCTTTGTTTTGCTTGCCCTCTGATTATCTG
TACTGTGGGTTTTAAGTATGCTACTTTCTCTCAGCATCCAATAATCATGGCCCTCAATTTATTTGTGGTCACCCA
GGGTTTCAAGCAAGAAGTCTTGCTTTATACAAATGTATCCATAAAATATCAGAGCTTGTTGGGCATGAACATCAA
ACTTTTGTTCCTAATATGGCTCTGTTTGGAAAAAAGTCAAAATCAGAAAGATGATTTGCAGAAAGAAAGAAA
AACTATGGTGTAATTTAAACTCTGGGCAGCCTCTGAATGAAATGCTACTTTCTTTAGAAATATAATAGCTGCCTT
AGACATTATGAGGTATACAACTAGTATTTAAGATACCATTTAATATGCCCCGTAAATGTCTTCAGTGTTCTTCAG
GGTAGTTGGGATCTCAAAAGATTTGGTTCAGATCCAAACAAATACACATTCTGTGTTTTAGCTCAGTGTTTTCTA
AAAAAAGAACTGCCACACAGCAAAAAATTGTTTACTTTGTTGGACAAACCAATCAGTTCTCAAAAAATGACCG
GTGCTTATAAAAAGTTATAAATATCGAGTAGCTCTAAACAAACACCTGACCAAGAGGGAAGTGAGCTTTGTGCT
TAGTATTTACATTGGATGCCAGTTTTGTAATCACTGACTTATGTGCAAACTGGTGCAGAAATCTATAAACTCTT
TGCTGTTTTTGATACCTGCTTTTTGTTTCATTTTGTGTTTTGTTTGTAAATGATAAACTTCAGAAAATAAAAT
GTCAGTGTTGAATAAAAAAAAAAAAAAAAAAAAAA

555/6881
FIGURE 518

MSKNTVSSARFRKVDVDEYDENKFVDEEDGGDQGAGPDEGEVDSCLRQGNMTAALQAALKNPPINTKSQAVKDRA
GSIVLKVLI SFKANDIEKAVQSLDKNGVDLLMKYIYKGFESPSDNSSAMLLQWHEKALAAGGVGSIVRVLTARKT
V

556/6881
FIGURE 519A

GCCAGGCAGAGTCCTACTATAGGCATGCAGCTCAGCTTGTCCCTCCAATGAAGCATTGCTGTGAAGTTCCTTT
CCCAGCTGCCTCCACTAATCTGCAAAAAGCACTTTCTAAAGCACTGGAAAGCCGAGATGAGGTGAAAACCAAGTG
GGGTGTTTTCTGACTTCATCAAGGCCCTTTATTAAATTCACGGTCATGTGTACCTGAGTAAGAGCTTGAAAAAGTT
GAGCCCTCTTCGAGAGAAAATTGGAAGAACAGTTTAAAGAGGCTGCTATTCCAAAAAGCTTTCAACTCTCAGCAGTT
AGTTCATGTCACTGTCTAATTAACCTGTTTCAACTTCATCACCTTCGTGACTTTAGCAATGAAACCGAGCAGCACAC
TTATAGCCAAGATGAGCAGCTATGTTGGACACAGTTGCTGGCCCTCTTTATGCTCTTTTCTTGGCATCCTGTGCAA
GTGTCTCTACAGAAATGAGTCTCAGGAGGAGTCTACAAATGCCTATCCTCTTCCAGCAGTCAAGGTCTCCATGGA
CTGGCTAAGACTCAGACCCAGGGTCTTTTCAAGAGGAGTGGTGGATGAAAGACAGTACATTGTCCTGCTGAT
TTCTCTTCTGAATAGTTTCCATCCCCATGAAGAGGACCTCTCAAGTATTAGTGCGACACCACTTCCAGAGGAGTT
TGAATTACAAGGATTTTGGCATTGAGACCTTCTTTTCAAGAACTTGGATTTTCCAAAGGTCACCAAGGTATTAC
AGGGGACAAAGAAGGCCAGCAACGACGAATACGACAGCAACGCTTGATCTCTATAGGCAAATGGATTGCTGATAA
TCAGCCAAGGCTGATTCACTGTGAAAATGAGGTAGGGAAATTTGTTGTTTATCACAGAAATCCCAGAATTAATACT
GGAAGACCCCAAGTGAAGCCAAAGAGAACCTCATTCTGCAAGAAACATCTGTGATAGAGTCGCTGGCTGCAGATGG
GAGCCCAGGGCTAAAATCAGTGCTATCTACAAGCCGAAATTTAAGCAACAACCTGTGACACAGGAGAGAAGCCAGT
GGTTACCTTCAAAGAAAACATTAAGACACGAGAAGTGAACAGAGACCAAGGAAGAAGTTTTCTCCCAAAGAGGT
GAGAAGGGACTATAGCAAAGGAATAACTGTAACATAAGATGATGGAAAGAAGGACAACAACAAGAGGAAAACCTGA
AACCAAGAAATGCACCTTAGAAAAGTTACAGGAAACAGGAAAGCAGAATGTGGCAGTGCAGGTAAAATCCCAGAC
AGAACTAAGAAAGACTCCAGTGTCTGAAGCCAGAAAAACACCTGTAACCTCAAACCCCAACTCAAGCAAGTAACTC
CCAGTTCATCCCCATTATCACCTTGGAGCCTTCCCTCCTCTTCCAGCAGGCCAGGGTTTCCGCCCCCAACATA
TGTTATCCCCCGCCTGTGGCATTCTTCTATGGGCTCAGGTTACACCTTCCAGCTGGTGTCTGTCTCCAGGAAC
CTTCTTTCAGCTACAGCTCACTCTCCAGCAGGAAACAGGTGCAAGCTGGGAAACAGTCCCACATTCTTTACAG
CCAGCAACGGCCCTCTGGACAGGGCCAAATGAACCAGGGACCTCAACAATCACAGCCACCTTCCAGCAACCCCT
TACATCTTTACCAGCTCAGCCAACAGCAGTCTACAAGCCAGCTGCAGGTTCAAGCTCTAACTCAGCAACAACA
ATCCCTTACAAAAGCTGTGCCGGCTTTGGGGAAAAGCCCGCTCACCACTCTGGATTCCAGCAGTATCAACAGGC
AGATGCCTCCAAACAGCTGTGGAATCCCCCTCAGGTTCAAGGCCCATTAGGGAAAATTATGCCTGTGAAACAGCC
CTACTACCTTCAGACCCAAGACCCCATAAAACCTGTTTGAAGCGTCATTGCAACCTCCTGTAATGCAGCAGCAGCC
TCTAGAAAAAAAATGAAGCCTTTTCCCATGGAGCCATATAACCATAATCCCTCAGAAGTCAAGGTCCCAGAATT
CTACTGGGATTCTTCTACAGCATGGCTGATAACAGATCTGTAATGGCACAGCAAGCAAACATAGACCGCAGGGG
CAAACGGTCACCAGGAATCTTCCGTCCAGAGCAGGATCTGTACCCAGAATGCCGTTTGAGAAATCCTTATTGGA
GAAGCCCTCAGAGCTCATGTACATTATCCTCTTCTGTCCCTCACCGGATTCTCTCTCAATCAGGAAAGATA
CCCAATAATAGTATGTTCAATGAGGTATATGGGAAAACCTGACATCCAGCTCCAAAGCAGAACTCAGTCCCTC
AATGGCCCCCAGGAAACATCTCTGTATTCCCTTTTTGAAGGGACTCCGTGGTCTCCATCACTTCCGTCCAGTTC
AGATCAATCAACACCAGCCAGCCAGTCTCCTCATTCTCTAACCAGCAGCCTACCCAGCTCTCCTCCAACACA
CAACCATAATTCTGTTCCATTCTCCAATTTTGGACCCATTGGGACTCCAGATAACAGGGATAGAAGGACTGCAGA
TCGGTGGAAAACCTGATAAGCCAGCCATGGGTGGGTTTGGCATTGATTATCTCTCAGCAACGTCATCCTCTGAGAG
CAGTTGGCATCAGGCCAGCACTCCGAGTGGCACCTGGACAGGCCATGGCCCTCCATGGAGGATTCTCTGTCTGT
CCTCATGGAAAGCCTAAAGTCTATCTGGTCCAGTTCCATGATGCATCCTGGACCTTCTGCTCTGGAGCAGCTGTT
AATGCAGCAGAAGCAGAAACAGCAACGGGGACAAGGCACCATGAACCTCCACACTGAGGCCAAAGTGGCAACCT
GGGAATGAAGGCTCCATAAACCATGGCATGTTGGGTTTGCAGGACTGGCCACACAGTCCCCTGCAGGTGGCAGC
CCTCTTTCTGTTTCTGCTGTCAAGAGGGTGTAAAGTATTCCACCAGCCCGCTGAGTGTGCACGAAATGTTGCA
GTGCAACAAAAAAGAAAAATCCATCAGGAACCTTCCGTCCCCCGGGCCCTTCCGAGGGAGAGAGAGAGGAACTG
CTGTTTATCTCTCAGTCAAGTTACTTGGTATCACCGCTCTCACCTTCTCCATCGTGCATGTCCCCAGCCACATGGGA
AGTGAAGCTGAGAAGGGAAGGCAGATGGGAGAAGCCAATGGGAACCTTCTCAGTCTTTTTTCTCTTTGGGGAA
TAAATAGGAATCCATTAATGATTGCTTTGCTGACTGAGAATGTAGTTGAAATTAACATCTTTTATTATTATTA
CTCTCAGTAGTAAATATCACACTGAATTCTTCCATACACAGGTGTGCTTCTAGTCAGTGTGTAGCAAGGAAAGC
CCCGTTCACTGCTCCTGTGAGAGGTTGGTGGTGACAGGATGGGGAACCGACCTCTTCAGCCAGTGGAAATGTTCC
ATAAGGGAGAGTTCAAGGCCGTGTCAGAAGGCTCTGGTAGGCCCTTCTCTGGCCAGGAGACTCCAGCAGGGAATGC
CCTTCACTCTGTAGGTGCTCGAGCCCCAATCGAGGATACAGTGTGGGTGGTGGTGTGGGCTGGACTAGCAGGTA

557/6881
FIGURE 519B

GACTGCTGTAGGATTTCAAATTACGTTTTTGTTCCTGTACATTTTACAGTCGCACAGCAAGCAGTCTCACAGAA
GGCAGGCTAGTCCATTACACAGCCTGACACGTTCTAATAGGTAGAAGCTTTTCAGTGTGGTTATTTTTCTTTGGTT
GGTTTTTGTGCCCCCATTCTACTTCCCACCCTCCTGCCCCATCTCCATCCCTTCTTTTACCCAATGCTGTATGCT
AGTAATTGTTTTTATTCTAATGTGTGCAACATCACATCTCCCCAAGAAGCAACAGCATGGGGTCCAGCAGTTGG
GGCCCAAAGACAGTCTGAAGAGGAAGGAAGCAGCAGTATCTGCGTAGCCACAGAGGGCCAGGCCCCTGCCCA
GCTGCAGTCTCCAGCCTCCACTTTCAGAGTGAATTTCAAGGCAGCACGGACATGTGCCCATCAGGCACAGAAGA
AAACACGACGTCGTCCATTTTGAAGAGACGAAAGAAAGGAAAATAAACTCTTTGTATGATATTTATTAGGAGGA
AAGAGGACTGAAAATGTTCTTGTGTAGAAACAGAAGGACAGCATTCTGTTAGTCATTTCTGGAAAAGTAATAT
TTTAAGGGGAAATTATGGAAACAATCTAATTGTTCAATTGCTGTGCTAGTGGTAGGGTTTATTTTCTGGGAGGTC
TCTCCTTTGTGTGCTGTATGTTTGTGTACACACACGTGCCCATCTGCTGTCCCAGAGGGGAGGGGTTGTGTGTG
CGAGTGTATGGAGTTAGTGTGGAACCTTAAGAGCTGGAAGACAGCTGTAGAGCAAAGCACATCCAGGAGCCCCAGT
TGTCACCTGCAGTCTGGGCAACCCAGCAATGAAAAGGGGTGAGATAACGCTCATTGCTCTTCAGAGAGAGTGGTT
GGAGCCCCCCCCCGCCCCGTATGCTTACATTATTGCTCTTTTAGTTTGACATGGTGTGTTGGGTTTTGTTTTTTGA
AAGGTCTGAAAAGGTGAAGCCCCCTACCCAATGGCAATATGAAACCTTTTGTGCTTCTCTTCAGCCCCCTTCCCTG
TGTCCACCTTTCTCTCTCTTCCCAAGCCTTTTTCTACTACCTTTACCCAGTTTGTGTGTTTGAGCTCTGCATT
CAGGCAGCTGCAACATTCCAGTGTGTTGAACTGTCACTGATTCTTGCGCCCTAGACAAGCTAACCAGGTTTACCAT
CTCACTCCCAGTAATACCCAGCTCCTATCTAAAGCCCCATTCTGCATGAGAATTTGGTGTGTTGGAATGTTTTCTG
ACTCTTGGGGCGGGATTCTCGCCTTATCATCCTCACTGTGGAGTAATGAGGGGGAGGAGAATCTTTATCAGAAA
CTGGTTTTGTGTAGTAACTTTCTTTTGTGGTTTTTTGTTTTGTTTTCTGGGTTTTGTTTTTGTTTTTGTCTGT
GCAAGACCTGCAGCTGCTGAAAATCAGCTTTGCCTTTAATTAAACCATGTTCTCTCC

558/6881
FIGURE 520

MSFLGILCKCPLQNESQEESSYNAYPLPAVKVSMDWLRLRPRVFQEAVVDERQYIWPWLISLLNSFHPHEEDLSSI
SATPLPEEFELQGFLALRPSFRNLDFSQGHQGITGDKEGQRRIRQQRLISIGKWIADNQPRLIQCENEVGKLLF
ITEIPELILEDPSEAKENLILQETSVIESLAADGSPGLKSVLSTSRNLSNNCDTGEKPVVTFKENIKTREVNRDQ
GRSFPPKEVRRDYSKGITVTKNDGKKDNNKRKTETKKCTLEKLQETGKQNVAVQVKSQTELKTPVSEARKTPVT
QTPTQASNSQFIPIHHPGAFPPPLPSRPGFPPPTYVIPPPVAFSMGSGYTFFAGVSVPGTFLQPTAHSPAGNQVQA
GKQSHIPYSQQRPSGPGPMNQGPQQSQPPSQQPLTSLPAQPTAQSTSQLQVQALTQQQQSPTKAVPALGKSPPHH
SGFQQYQQADASKQLWNPPQVQGFLGKIMPVKQPYLQTDPIKLFEPQLQPPVMQQQPLEKKMKPFPMEPYNHN
PSEVKVPEFYWDSSYSMADNRSVMAQQANIDRRGKRSPGIFRPEQDPVPRMPFEKSLLEKPSSELMSSSSFLSLT
GFSLNQERYPNNSMFNEVYGNLTSSSKAELSPSMAPQETSLSLFEGTPWSPSLPASSDHSTPASQSPHSSNPS
SLPSSPPTHNNHNSVPFSNFGPIGTPDNRRRTADRWKTDKPAMGGFGIDYLSATSSSESSWHQASTPSGTWTGHG
PSMEDSSAVLMESLKSIWSSSMHFGPSALEQLLMQQKQKQQRGQGTMNPPH

559/6881

FIGURE 521A

ACCACCTGATCAAGGAAAAGGAAGGCACAGCGGAGCGCAGAGTGAGAACCACCAACCGAGGCGCCGGGCAGCGAC
CCCTGCAGCGGAGACAGAGACTGAGCGGCCCGGCCCGCCATGCCTGCGCTCTGGCTGGGCTGCTGCCTCTGCTT
CTCGCTCCTCCTGCCCGCAGCCCGGGCCACCTCCAGGAGGGAAGTCTGTGATTGCAATGGGAAGTCCAGGCAGTG
TATCTTTGATCGGGAACCTCACAGACAAACTGGTAATGGATTCCGCTGCCTCAACTGCAATGACAACACTGATGG
CATTCACTGCGAGAAGTGCAAGAATGGCTTTTACCGGCACAGAGAAAGGGACCGCTGTTTGCCCTGCAATTGTAA
CTCCAAAGGTTCTCTTAGTGCTCGATGTGACAACCTCCGGACGGTGACGCTGTAAACCAGGTGTGACAGGAGCCAG
ATGCGACCGATGTCTGCCAGGCTTCCACATGCTCACGGATGCGGGGTGCACCCAAGACCAGAGACTGCTAGACTC
CAAGTGTGACTGTGACCCAGCTGGCATCGCAGGGCCCTGTGACGCGGGCCGCTGTGTCTGCAAGCCAGCTGTCAC
TGGAGAACGCTGTGATAGGTGTGATCAGGTTACTATAATCTGGATGGGGGGAACCCCTGAGGGCTGTACCCAGTG
TTTCTGTATGGGCATTACGCCAGCTGCCGACGCTCTGCAGAATACAGTGTCCATAAGATCACCTCTACCTTTCA
TCAAGATGTTGATGGCTGGAAGGCTGTCCAACGAAATGGGTCTCCTGCAAAGCTCCAATGGTCACAGCGCCATCA
AGATGTGTTTAGCTCAGCCCAACGACTAGACCTGTCTATTTTGTGGCTCCTGCCAAATTTCTTGGGAATCAACA
GGTGAGCTATGGTCAAAGCCTGTCTTTGACTACCGTGTGGACAGAGGAGGCAGACACCCATCTGCCATGATGT
GATTCTGGAAGGTGCTGGTCTACGGATCACAGCTCCCTTGATGCCACTTGGCAAGACACTGCCTTGTGGGCTCAC
CAAGACTTACACATTACAGTTAAATGAGCATCCAAGCAATAATTGGAGCCCCAGCTGAGTTACTTTGAGTATCG
AAGGTTACTGCGGAATCTCACAGCCCTCCGCATCCGAGCTACATATGGAGAATACAGTACTGGGTACATTGACAA
TGTGACCTGATTTTCAAGCCCGCCCTGTCTCTGGAGCCCCAGCACCCCTGGGTTGAACAGTGTATATGTCTGTGG
GTACAAGGGGCAATTTCTGCCAGGATTGTGCTTCTGGCTACAAGAGAGATTACAGCGAGACTGGGGCCTTTTGGCAC
CTGTATTCTTTGTAAGTGTCAAGGGGGAGGGGCCCTGTGATCCAGACACAGGAGATTGTTATTACAGGGGATGAGAA
TCCTGACATTGAGTGTGCTGACTGCCCAATTGGTTTCTACAACGATCCGCACGACCCCCGAGCTGCAAGCCATG
TCCCTGTCTATAACGGGTTCAAGTGTCTAGTGATGCCGAGACGGAGGAGGTGGTGTGCAATAACTGCCCTCCCGG
GGTCACCGGTGCCCGCTGTGAGCTCTGTGCTGATGGCTACTTTGGGGACCCCTTTGGTGAACATGGCCCAGTGAG
GCCTTGTGAGCCCTGTCAATGCAACAACAATGTGGACCCCAAGTGCCCTCTGGGAATTGTGACCGGCTGACAGGCAG
GTGTTTGAAGTGTATCCACAACACAGCCGGCATCTACTGCGACCAAGTGCAGAGCAGGCTACTTCGGGGACCCATT
GGCTCCCAACCCAGCAGACAAGTGTGAGCTTGCACACTGTAACCCCATGGGCTCAGAGCCTGTAGGATGTGCAAG
TGATGGCACCTGTGTTTGCAAGCCAGGATTGGTGGCCCCAACTGTGAGCATGGAGCATTACGCTGTCCAGCTTG
CTATAATCAAGTGAAGATTACAGTGGATCAGTTTATGCAGCAGCTTACAGAGAATGGAGGCCCTGATTTCAAAGGC
TCAGGGTGGTGTGAGTAGTACCTGATACAGAGCTGGAAGGCAGGATGCAGCAGGCTGAGCAGGCCCTTCAGGA
CATTCTGAGAGATGCCAGATTTCAAGAGGTGTAGCAGATCCCTTGGTCTCCAGTTGGCCAAGGTGAGGAGCCA
AGAGAACAGCTACCAGAGCCGCTGGATGACCTCAAGATGACTGTGGAAAGAGTTCGGGCTCTGGGAAGTCAGTA
CCAGAACCAGGTTCCGGGATACTCACAGGCTCATCACTCAGATGCAGCTGAGCCTGGCAGAAAGTGAAGCTTCTTT
GGGAAACACTAATCTTCTGCCCTCAGACCACTACGTGGGGCCAAATGGCTTTAAAAGTCTGGCTCAGGAGGCCAC
AAGATTAGCAGAAAGCCACGTTGAGTCAGCCAGTAACATGGAGCAACTGACAAGGGAACTGAGGACTATTCCAA
ACAAGCCCTCTCACTGGTGCAGCAAGGCCCTGCATGAAGGAGTCGGAAGCGGAAGCGGTAGCCCGGACGGTGCTGT
GGTGAAGGGCTTGTGGAAAAATTTGGAGAAAACCAAGTCCCTGGCCCAGCAGTTGACAAGGGAGGCCACTCAAGC
GGAAATTGAAGCAGATAGGTCTTATCAGCACAGTCTCCGCCTCCTGGATTACGTGTCTCGGCTTCAGGGAGTCAG
TGATCAGTCCTTTCAGGTGGAAGAAGCAAAGAGGATCAAACAAAAGCGGATTCACCTCAAGCCTGGTAACCAAG
GCATATGGATGAGTTCAAGCGTACACAGAAGAATCTGGGAACTGGAAAGAAGAAGCACAGCAGCTCTTACAGAA
TGGAAAAAGTGGGAGAGAGAAATCAGATCAGCTGCTTTCCCGTGCCAATCTTGCTAAAAGCAGAGCACAGAAGC
ACTGAGTATGGGCAATGCCACTTTTATGAAGTTGAGAGCATCCTTAAAAACCTCAGAGAGTTTGACCTGCAGGT
GGACAACAGAAAAGCAGAAGCTGAAGAAGCCATGAAGAGACTCTCCTACATCAGCCAGAAGGTTTCAGATGCCAG
TGACAAGACCCAGCAAGCAGAAAAGAGCCCTGGGGAGCGCTGCTGTGATGCACAGAGGGCAAAGAATGGGGCCGG
GGAGGCCCTGGAAATCTCCAGTGAGATTGAACAGGAGATTGGGAGTCTGAACTTGAAGCCAATGTGACAGCAGA
TGGAGCCTTGGCCATGGAAAAGGGACTGGCCTCTCTGAAGAGTGAAGTGAAGGGAAGTGAAGGAGAGCTGGAAG
GAAGGAGCTGGAGTTTGACACGAATATGGATGCAGTACAGATGGTGATTACAGAAGCCCAAGAGGTTGATACCAG
AGCCAAGAACGCTGGGGTTACAATCCAAGACACACTCAACACATTAGACGGCCTCCTGCATCTGATGGACCAGCC
TCTCAGTGTAGATGAAGAGGGGCTGGTCTTACTGGAGCAGAAGCTTTCCCGAGCCAAGACCCAGATCAACAGCCA
ACTGCGGCCCATGATGTGAGAGCTGGAAGAGAGGGCACGTGAGCAGAGGGGCCACCTCCATTTGCTGGAGACAAG

560/6881
FIGURE 521B

CATAGATGGGATTCTGGCTGATGTGAAGAACTTGGAGAACATTAGGGACAACCTGCCCCAGGCTGCTACAATAC
CCAGGCTCTTGAGCAACAGTGAAGCTGCCATAAATATTTCTCAACTGAGGTTCTTGGGATACAGATCTCAGGGCT
CGGGAGCCATGTCATGTGAGTGGGTGGGATGGGGACATTTGAACATGTTTAAATGGGTATGCTCAGGTCAACTGAC
CTGACCCCATTCCTGATCCCATGGCCAGGTGGTTGTCTTATTGCACCATACTCCTTGCTTCCTGATGCTGGGCAA
TGAGGCAGATAGCACTGGGTGTGAGAATGATCAAGGATCTGGACCCCAAAGAATAGACTGGATGGAAAGACAAAC
TGCACAGGCAGATGTTTGCCTCATAATAGTCGTAAGTGGAGTCCCTGGAATTTGGACAAGTGCTGTTGGGATATAG
TCAACTTATTCTTTGAGTAATGTGACTAAAGGAAAAAACTTTGACTTTGCCCAGGCATGAAATTCTTCCTAATGT
CAGAACAGAGTGCAACCCAGTCACACTGTGGCCAGTAAAATACTATTGCCTCATATTGTCCTCTGCAAGCTTCTT
GCTGATCAGAGTTCCTCCTACTTACAACCCAGGGTGTGAACATGTTCTCCATTTTCAAGCTGGAAGAAGTGAGCA
GTGTTGGAGTGAGGACCTGTAAGGCAGGCCCATTCAGAGCTATGGTGCTTGCTGGTGCCCTGCCACCTTCAAGTTC
TGGACCTGGGCATGACATCCTTTCTTTAATGATGCCATGGCAACTTAGAGATTGCATTTTATTAAAGCATTTTC
CTACCAGCAAAGCAAATGTTGGGAAAGTATTTACTTTTTCGGTTTCAAAGTGATAGAAAAGTGTGGCTTGGGCAT
TGAAAGAGGTAAAATTCTCTAGATTATTAGTCCTAATTCAATCCTACTTTTTCGAACACCAAAAATGATGCGCAT
CAATGTATTTTATCTTATTTTCTCAATCTCCTCTCTCTTTTCTCCACCCATAATAAGAGAATGTTCTACTCACA
CTTCAGCTGGGTACATCCATCCCTCCATTATCCTTCCATCCATCTTTCCATCCATTACCTCCATCCATCCTTC
CAACATATATTTATTGAGTACCTACTGTGTGCCAGGGGCTGGTGGGACAGTGGTGACATAGTCTCTGCCCTCATA
GAGTTGATTGTCTAGTGAGGAAGACAAGCATTTTTAAAAAATAAATTTAAACTTACAACTTTGTTTGTACAAAG
TGGTGTATTTATTGCAATAACCGCTTGGTTTGCAACCTCTTTGCTCAACAGAACATATGTTGCAAGACCCTCCCATG
GGGGCACTTGAGTTTGGCAAGGCTGACAGAGCTCTGGGTGTGCACATTCTTTGCATTCCAGCTGTCACTCTG
TGCTTTTCTACAACTGATTGCAACAGACTGTTGAGTTATGATAACACCAGTGGGAATTGCTGGAGGAACCAGAGG
CACTTCCACCTTGGCTGGGAAGACTATGGTGCTGCCTTGCTTCTGTATTTCTTGGATTTTCTGAAAGTGT
TAAATAAAGAACAATTGTTAGA

561/6881
FIGURE 522

CCCAACCGCAGTTGACTAGCACCTGCTACCGCGCCTTTGCTTCCTGGCGCACGCGGAGCCTCCTGGAGCCTGCCA
CCATCCTGCCTACTACGTGCTGCCCTGCGCCCGCAGCCATGTGCCGCACCCTGGCCGCCTTCCCCACCACCTGCC
TGGAGAGAGCCAAAGAGTTCAAGACACGTCTGGGGATCTTTCTTCACAAATCAGAGCTGGGCTGCGATACTGGGA
GTACTGGCAAGTTCGAGTGGGGCAGTAAACACAGCAAGAGAATAGAACTTCTCAGAAGATGTGCTGGGGTGA
GAGAGTCGTTTCGACCTGCTGCTGAGCAGTAAAAATGGAGTGGCTGCCTTCCACGCTTTCCTGAAGACAGAGTTCA
GTGAGGAGAACCTGGAGTTCTGGCTGGCCTGTGAGGAGTTCAAGAAGATCCGATCAGCTACCAAGCTGGCCTCCA
GGGCACACCAGATCTTTGAGGAGTTCAATTGCAGTGAGGCCCTAAAGAGGTCAACATTGACCATGAGACCCAGG
AGCTGACGAGGATGAACCTGCAGACTGCCACAGCCACATGCTTTGATGCGGCTCAGGGGAAGACACGTACCCTGA
TGGAGAAGGACTCCTACCCACGCTTCTGAAGTCGCTTACCGGGACCTGGCTGCCCAAGCCTCAGCCGCCT
CTGCCACTCTGTCCAGCTGCAGCCTGGACGAGCCCTCACACACCTGAGTCTCCACGGCAGTGAGGAAGCCAGCCG
GGAAGAGAGGTTGAGTCAACCATCCCCGAGGTGGCTGCCCTGTGTGGGAGGCAGGTTCTGCAAAGCAAGTGCAA
GAGGACAAAAAAGAGTCAACCATCCCCGAGGTGGCTGCCCTGTGTGGGAGGCAGGTTCTGCAAAGCAAGTGCAA
GATACTGTGGGACTCATGCTGGAGAGGAGCCGCCACTTCCAGGACCTGTGAATAAGGGCTAATGATGAGGGTTG
GTGGGGCTCTCTGTGGGGCAAAAGGTGGTATGGGGGTTAGCACTGGCTCTCGTTCTCACCAGGAGAAGGAAGTGT
TCTAGTGTGGTTTATAGAAACATGTGGATAAAGGGAACCATGAAAATGAGAGGAGGAAAGACATCCAGATCAGCTG
TTTTGCCTGTGCTCAGTTGACTCTGATTGCATCCTGTTTTCTTAATTCACAGACTGTTCTGGGCACGGAAGGGA
CCCTGGATGTGGAGTCTTCCCTTTGGCCCTCCTCACTGGCCTCTGGGCTAGCCAGAGTCCCTTAGCTTGTACC
TCGTAACACTCCTGTGTGTCTGTCCAGCCTTGCAAGTCAAGGCCAGCAAGCTGATGTGACTCTTGCCCAT
GCGAGATATTTATACCTCAAACACTGGCCTGTGAGCCCTTTCCAAGTCAAGTGGAGAGCCCTGAAAGGAGGCTCAC
TTGAATCCAGCTCAGTGCTCTGGGTGGCCCCCTGCAGGTGGCCCCCTGACCCTGCGTTGCAGCAGGGTCCACCTGT
GAGCAGGCCCGCCCTGGGGCCTCTTCTGGATGTGCCCTCTCTGAGTCTGTGCTGTCTCTTGGAGGCAGGGCCC
AGGAGAACAAAGTGTGGAGGCTCGGGGAGTGGCTTTTCCAGCTCTCATGCCCCGAGTGTGGAACAAGGCAGAA
AAGGATCCTAGGAAATAAGTCTCTTGGCGGTCCCTGAGAGTCTGTGTAATCCAGCCAGTGTTTTTTGTGGTAT
GAGAACAGGCAAAAGAGATGCCCCGAGATAGAAGGGGAGCCTTGTGTTTCTTCTGCAGACGTGAGATGAACA
CTGGAGTGGGCAGAGGTGGCCAGGACCATGGCACCTTAGAGTGCAGAAGCTGGGGGAGAGGCTGCTTGAAG
GGCAGGACTGGGGATAATCAGAACCTGCCTGTACCTCAGGGCATCACTGAACAAACATTTCTGATGGCAACTC
CTGCGGCAGAGCCAGGCTGGGGAAGTGAACATCCAGGGCAGCCCTTTGTGGCCAGGATAATCAACACTGTT
CTCTCTGTACCATGAGCTCCTCCAGGAGATTATTTAAGTGTATTGTATCATTGGTTTTCTGTGATTGTCATAACA
TTGTTTTTGTATTGTTGGTGTCTGTTATTTATTATTGTAATTTAGTTTGCCTCTACTGGAGAATCTCAGCA
GGGTTTTAGCCTGACTGTCTCCCTTTCTCTACCAGACTCTACCTCTGAATGTGCTGGGAACCTCTTGGAGCCTG
TCAGGAACCTCCTCACTGTTTAAATATTTATTATTGTGACAAATGGAGCTGGTTTTCTAGATATGAATGATGTTT
GCAATCCCCATTTTCTGTTTTAGCATGTTATATTCTTATAAAATAAAAGCAAAAGTCAAATATG

562/6881
FIGURE 523

GGGTGCGGGGCTGCTGGCGGCTCTGCAGAGTCGAGAGTGGGAGAAGAGCGGAGCGTGTGAGCAGTACTGCGGCCT
CCTCTCCTCTCCTAACCTCGCTCTCGCGGCTAGCTTTACCCGCCCCGCTGCTCGGCGACCAGCGGGGATCCTCC
CCAGCCGCAAGTCCACGAAGAAAGCAACGAATGAAAATTATGAAGACAACGAGAAGTCAGACTCCTCCGGGTGCG
CGTCCAGCTGCTTCGGCTTCGTCGCCTACTCTGTGAACTCCGGGGAGAGATCTCGAGTCAAGATTAAGACCTTA
ACCCACCAACCTGCCTGTTCGGACACCCCCGGGCGGCCGCTGTCTGTCCCCTTCTCCATCGCCCTCTCCCAGA
AAGCTCCGGTGTCTGGACCAGCTAGAGTCTGAGAAAGAGGAGAGGCGCGAACGCCACTCCAAAAAGAGAAGGGTT
AAAGAGGGCAACCCTAACGATACGCTTGACTTTCTGTGGCTGGGAACACCTTCCACCATGACCACCTCAGCAAGT
TCCCACTTAAATAAAGGCATCAAGCAGGTGTACATGTCCCTGCCTCAGGGTGAGAAAGTCCAGGCCATGTATATC
TGGATCGATGGTACTGGAGAAGGACTGCGCTGCAAGACCCGGACCCCTGGACAGTGAGCCCAAGTGTGTGGAAGAG
TTGCCTGAGTGGAAATTCGATGGCTCTAGTACTTTACAGTCTGAGGGTTCCAACAGTGACATGTATCTCGTGCCT
GCTGCCATGTTTCGGGACCCCTTCCGTAAGGACCCTAACAGCTGGTGTATGTGAAGTTTCAAGTACAATCGA
AGGCCTGCAGAGACCAATTTGAGGCACACCTGTAAACGGATAATGGACATGGTGAGCAACCAGCACCCCTGGTTT
GGCATGGAGCAGGAGTATACCCCTCATGGGGACAGATGGGCACCCCTTTGGTTGGCCTTCCAACGGCTTCCCAGGG
CCCCAGGGTCCATATTACTGTGGTGTGGGAGCAGACAGAGCCTATGGCAGGGACATCGTGGAGGCCATTACCGG
GCCTGCTTGTATGCTGGAGTCAAGATTGCGGGGACTAATGCCGAGGTCTGCTGCCAGTGGGAATTCAGATT
GGACCTTGTGAAGGAATCAGCATGGGAGATCATCTCTGGGTGGCCCGTTTCATCTTGCATCGTGTGTGTGAAGAC
TTTGAGTGTATAGCAACCTTTGATCCTAAGCCCATTCCTGGGAATGGAATGGTGCAGGCTGCCATACCAACTTC
AGCACCAAGGCCATGCGGGAGGAGAATGGTCTGAAGTACATCGAGGAGGCCATGAGAAACTAAGCAAGCGGCAC
CAGTACCACATCCGTGCCTATGATCCCAAGGGAGGCCTGGACAATGCCCGACGTCTAACTGGATTCCATGAACCC
TCCAACATCAACGACTTTTCTGCTGGTGTAGCCAATCGTAGCGCCAGCATACGCATTCCCCGGACTGTTGGCCAG
GAGAAGAAGGGTTACTTTGAAGATCGTCGCCCCCTCTGCCAACTGCGACCCCTTTTCGGTGACAGAAGCCCTCATC
CGCACGTGTCTTCTCAATGAAACCGCGATGAGCCCTTCCAGTACAAAATTAAGTGGACTAGACCTCCAGCTGT
TGAGCCCTCCTAGTTCTTCATCCCACTCCAACCTTCCCCCTCTCCAGTTGTCCCGATTGTCCGATTGAACAGGGT
GGAATATCAAGGTCGTTTTTTTTCATTCCATGTGCCAGTTAACTTGTCTTTCTTTGTTTGGCTGGGATAGAGGGG
TCAAGTTATTAATTTCTTCACACCTACCCTCCTTTTTTTTCCCTATCACTGAAGCTTTTTAGTGCATTAGTGGGGA
GGAGGGTGGGGAGACATAACCACTGCTTCCATTTAATGGGGTGACCTGTCCAATAGGCGTAGCTATCCGGACAG
AGCAGTTCGAGAAAGGGGCTCTCTTCTCCAGGTAGCTGAAAGGGGAAGACCTGACGTACTCTGGTTAGGTTAG
GACTTGCCCTCGTGGTGGAACTTTTCTTAAAAAGTTATAACCAACTTTTCTATTAAAAAGTGGGAATTAGGAGAG
AAGGTAGGGGTGGGAATCAGAGAGAATGGCTTTGGTCTCTTGTCTTGTGGGACTAGCCTGGCTTGGGACTAAATG
CCCTGCTCTGAACACGAAGCTTAGTATAAACTGATGGATATCCCTACCTTGAAAGAAGAAAAGGTTCTTACTGCT
TGGTCCTTGATTTATCACACAAAGCAGAAATAGTATTTTTATATTTAAATGTAAAGACAAAAACTATATGTATGG
TTTTGTGATTATGTGTGTTTTGCTAAAGGAAAAAACCATCCAGGTACAGGGGCACCAAATTTGAGACAAATAGT
CGGATTAGAAAATAAAGCATCTCATTTTGTAGTAGAGAGCAAGGGAAGTGGTTCTTAGATGGTGATCTGGGATTAGG
CCCTCAAGACCCCTTTTGGGTTTCTGCCCTGCCACCCCTCTGGAGAAGGTGGGCACTGGATTAGTTAACAGACAAC
ACGTTACTAGCAGTCACCTGATCTCCGTGGCTTTGGTTTTAAAGACACACTTGTCCACATAGGTTTAGAGATAAG
AGTTGGCTGGTCAACTTGAGCATGTTACTGACAGAGGGGTATTGGGGTTATTTTCTGGTAGGAATAGCATGTCA
CTAAAGCAGGCCCTTTTGATATTAAATTTTTTAAAAAGCAAAATTATAGAAGTTTAGATTTTAATCAAATTTGTAG
GGTTTCTAGGTAATTTTTACAGAATTGCTTGTGTTGCTTCAACTGTCTCCTACCTCTGCTCTTGGAGGAGATGGGG
ACAGGGCTGGAGTCAAAACACTTGTAATTTTGTATCTTGATGTCTTTGTTAAGACTGCTGAAGAATTATTTTTTT
TCTTTTATAATAAGGAATAAACCCACCTTTATTCCTTCATTTTCATCTACCATTTTCTGGTTCTGTGTGGCTG
TGGCAGGCCAGCTGTGGTTTTCTTTTGCCATGACAACTTCTAATTGCCATGTACAGTATGTTCAAAGTCAAATAA
CTCCTCATTGTAAACAACTGTGTAAGTGGCCAAAGCAGCACTTATAAATCAGCCTAACAT

563/6881
FIGURE 524A

GC GGGGAGGGGCGGGGCTATGGAGAGGAGGAGGAAGATGGCGGGCGGGCTGCTCTGAAGAGACCTCGGCGGGCGG
GGAGGAGGAGAGAAGCGCAGCGCCGCGCCGCGCCGGGGGCCATGTGGGAGGAGTCGGAGTCGCTGTTGCCGCCG
CCGCCTGTAGCTGCTGGACCCGAGTGGGAGTGAGGGGGAAACGGCAGGATGAAGTTCGCCGAGCACCTCTCCGCG
CACATCACTCCCGAGTGGAGGAAGCAATACATCCAGTATGAGGCTTTCAAGGATATGCTGTATTACAGTCAGGAC
CAGGCACCTTCTGTGGAAGTTACAGATGAGGACACAGTAAAGAGGTATTTTGCCAAGTTTGAAGAGAAGTTTTTC
CAAACCTGTGAAAAAGAACCTGCCAAAATCAACACATTTTATTAGAGAAGCTCGCAGAGGCTCAGCGCAGGTTT
GCTACACTTCAGAATGAGCTTCAGTCATCACTGGATGCACAGAAAGAAAGCACTGGTGTTACTACGCTGCGACAA
CGCAGAAAGCCAGTCTTCCACTTGTCCCATGAGGAACGTGTCCAACATAGAAATATTAAAGACCTTAAACTGGCC
TTCAGTGAGTTCTACCTCAGTCTAATCCTGCTGCAGAACTATCAGAATCTGAATTTTACAGGGTTTCGAAAAATC
CTGAAAAAGCATGACAAGATCCTGGAAACATCTCGTGAGCAGATTGGCGAGTGGCTCAGTAGAGGTGGCCCCA
TTTTATACATGCAAGAAAATCAACCAGCTTATCTCTGAAACTGAGGCTGTAGTGACCAATGAACCTGAAGATGGT
GACAGACAAAAGGCTATGAAGCCTTTACGTGTCCCCCTTTGGGAGCTGCTCAGCCTGCACCAGCATGGACTACT
TTTAGAGTTGGCTATTTTGTGGAATATTCATTGTACTGAATATTACCCTTGTGCTTGGCGCTGTATTTAACTT
GAAACAGATAGAAGTATATGGCCCTTGATAAGAATCTATCGGGGTGGCTTTCTTCTGATTGAATTCCTTTTTCTA
CTGGGCATCAACAGTATGGTTGGAGACAGGCTGGAGTAAACCATGTACTCATCTTTGAACTTAATCCGAGAAGC
AATTTGTCTCATCAACATCTCTTTGAGATTGCTGGATTCTCGGGATATTGTGGTGCCTGAGCCTTCTGGCATGC
TTCTTTGTCTCCAATTAGTGTATCCCCACATATGTGTATCCACTTGCCTTTATGGATTATGGTTTTCTTCCTT
ATCAACCCACCAAACTTTTCTACTATAAATCCCGGTTTTGGCTGCTTAACTGCTGTTTCGAGTATTTACAGCC
CCCTTCCATAAGGTAGGCTTTGTGATTTCTGGCTGGCGGATCAGCTGAACAGCCTGTCACTGATAGTGGAC
CTGGAATATATGATCTGCTTCTACAGTTTGGAGCTCAAATGGGATGAAAGTAAGGGCCTGTGCCAAATAATTCA
GAAGAATCAGGAATTTGCCACAAATATACATATGGTGTGCGGGCCATTGTTCACTGCTTCTGCTTGGCTTCGC
TTCATCCAGTGCCTGCGCCGATATCGAGACACAAAAGGGCCTTTCTCATTTAGTTAATGCTGGCAAATACTCC
ACAACCTTTCTTATGGTGACGTTTGCAGCCCTTTACAGCACTCACAAAGAACGAGGTCAGTGGACACTATGGTG
TTCTTTTACCTGTGGATTGTCTTTTATATCATCAGTTCTGCTATACCCTCATCTGGGATCTCAAGATGGATGG
GGTCTCTTCGATAAGAAATGCTGGAGAGAACACTTTCTCCGGGAAGAGATTGTATACCCCCAAAAGCCTACTAC
TACTGTGCCATAATAGAGGATGTGATTCTGCGCTTTGCTTGGACTATCCAAATCTCGATTACCTCTACAACCTTG
TTGCCTCATTCGCGGACATCATTGCTACTGTCTTTGCCCACTTGAGGTTTTCCGGCGATTGTGTGGAACCTTC
TTCCGCCTGGAGAATGAACATCTGAATAACTGTGGTGAATTCGCTGTGTGCGGGACATCTCTGTGCCCCCTG
AACGCAGATGATCAGACTCTCCTAGAACAGATGATGGACCAGGATGATGGGTACGAAACCGCCAGAAGATCGG
TCATGGAAGTACAACCAGAGCATATCCCTGCGCCGGCCTCGCTCTCAATCCAAGGCTCGTGACACTAAG
GTATTGATAGAAGACACAGATGATGAAGCTAACACTTGAATTTCTGAAGCTAGCTTAAACATCTTTGGTTTTCC
TACTCTACAATCCTTTCTCGACCAACGCAACCTCTAGTACCTTTCCAGCCGAAAACAGGAGAAAACACATAACA
CATTTTCCGAGCTCTTCCGGATCGGATCCTATGGACTCCAAACAAGCTCACTGTGTTTTCTTTCTTTCTTCTGG
TTTAATTTTAATTTTCTATTTTCAAAACAAATATTTACTTCATTGCCAATCAGAGGATGTTTTAAGAAACAAA
CATAGTATCTTATGGATTGTTTACAATCACAAGGACATAGATACCTATCAGGATGAAGAACAGGCATTGCAAGGA
CCCTCTGATGGGACGGTACTGAGATATCTCGGCTTCCGCTCAGCCCGTTTTGACTGGTTGAAACCGGACATTGG
TTTTTAAATTTTTTGTGCTTATGTGGAGAATTTTTTCTTTCTTCTCATACCCAGCGCAAAGGCACTGGCCGCA
CTTGCAGGAAAAGTGCAACTTAAAGCAGTACCTTCATTCAATGAAGCTACTTTTTAATTTGATGTAACCTTTCTTA
TTTTGGGAAGGGTGTCTGGGTGGGTGGGAAATATGATGTATTTGTTACACATAGTTTTCTCATTATTTATGAAAC
TTAACCATACAGAATGATATAACTCCTGTGCAATGAAGGTGATAACAGTAAAGAAAGGCAGGGGAAACTTACGTT
GGATGACATTTATGAGGTCAGTCCACATACCTCTTTCAAGTTGCTGCTGCTTTAGAGTCTGAGCATATCT
TTTGTTTTTATTTTAAGCCAAAGTTTCATTGCTAACTTCTTAAGTTGCTGCTGCTTTAGAGTCTGAGCATATCT
CTCATAACAAGGAATCCCACACTTCACACCACCGGCTGAATTTTATGGAAGAGGTTCTGATAATTTTTTTAACTT
TTTAAGGAACAGATGTGGAATACACTGGCCCATATTTCAACCTTAACAGCTGAAGCTATGCCTTATTATGCATCC
ACATGTATGGTCCCTGTAGCGTGACCTTTACTAGCTCTGAATCAGAAGACAGAGCTATTTAGAGGCTCTGTGTG
CCCTCACTAGATAGTTTTTCTTCTGGGTTCAACCACTTTAGCCAGAATTTGATCAAATTTAAAGTCTGTATGGG
GAAACTATATTTTGTAGCACATGGAACAAATTATACTTCTCATTATATGTTGATACAAAAGACCTTGGCA
GCCATTTCTCCAGCAGTTTTTAAAGGATGAACATTGGATTTTATGCCATCCCATAGAAAACCTGTTTTAAATTT

564/6881

FIGURE 524B

TAGGGATCTTTACTTGGTCATACATGAAAAGTACACTGCTTAGAAATTATAGACTATTATGATCTGTCCACAGTG
CCCATTGTCACITCTTTGTCTCATTCTTCCCTTTGTTCCCTTAGTCATCCAAATAAGCCTGAAAACCATAAGAGA
TATTACTTTATTGAATATGGTTGGCATTAAATTTAGCATTTCATTATCTAACAAAATTAATATAAATTCCAGGAC
ATGGTAAATGTGTTTTAATAACCCCCAGACCCAAATGAAAATTTCAAAGTCAATACCAGCAGATTTCATGAAAGT
AAATTTAGTCCTATAATTTTCAGCTTAATTATAAACAAAGGAACAAATAAGTGGAGGGCAGCTATTACCATTTCG
CTTAGTCAAAACATTTCGGTTACTGCCCTTTAATACACTCCTATCATCAGCACTTCCACCATGTATTACAAGTCTT
GACCCATCCCTGTCGTAAGTCCAGTAAAAGTTACTGTTACTAGAAAATTTTTATCAATTAAGTACAAAATAGTTT
CTTTTAAAGTAGTTTCTTCCATCTTTATTCTGACTAGCTTCCAAAATGTGTTCCCTTTTTGAATCGAGGTTTTT
TTGTTTTGTTTTGTTTTCTGAAAAAATCATACAACTTTTGTCCTTCTATTGCTTTTTTGTTGTTTTGTTAAGCATGT
CCCTTGCCCCAAATGGAAGAGGAAATGTTTAATTAATGCTTTTAGTTTAAATAAATTGAATCATTATAATAAT
CAGTGTTAACAATTTAGTGACCCTTGGTAGGTTAAAGGTTGCATTATTTATACTTGAGATTTTTTCCCCTAACT
ATTCTGTTTTTTGTACITTTAAACTATGGGGGAAATATCACTGGTCTGTCAAGAAACAGCAGTAATTATTACTGA
GTTAAATTGAAAAGTCCAGTGGACCAGGCATTTCTTATATAAATAAAATTGGTGGTACTAATGTG

565/6881
FIGURE 525

AAATAAATTATTTTTGTTAAGGCAAGCATTGGTGTGTTCTTTAACTTGCTACTTGGAGACCTAGTGTCCAGTCTGG
ACAGAAATGCACAGAGACCAGCCTCACCTGGAATGCAGCCAGACTCGATGTCCCTCAGCACACACAGCTCCTGGCC
ACAGACTGCCCATAGAACTGTCTGCACCCAAACCCCATGGCCTTTTCATGCACGGAGACAGGCCCTCTGGATGTGC
AGCCTTGCCACCCCTGCCCCAATCCTCCCTGAGAGCTCCTGCCTCAGTGCCCTGGGCTGGTGAGGGAGAAGCCT
GTCTGCACCTGCCTAATTCCAGCTCCTCCAGGAAGCAAGGGCTTGTTTGTGAGAGCTGCAGGGTTGGCCACTCGG
TGGGGAGGAGTCAGCCAGGATTAGAGAGCCTGCCCCCTAATCCGGCCTGCTGGGTTTTACAAGGATCAGAGCTGCT
GATAATGAACCTCATTAAAGGGGGAGCAGGAGCCTCAATCCGATTGGTTTTCTCTTTGACATCTTCACTCTGCTC
AGATGGCCTGGGTGCTATGTGGAGCAGGTGGGATGCCAAGGCCACTCCTGCTATGGGGCAGCTGGGGCTGGGGAG
GGATGGCAGTCTCCCTGCATGTTTTCCCTCGACCTCTTTAGCTGCAGCGCCTTGCTGGGCTCCTGGGTTGGACTCC
CTCTCTGTGCCCCCTGCTCCAGGCACCCATTGGCTCCATCCTCCTGGTTGTGCTCTGCACCCCTGCTCCCTTGGG
CTGGCCCTGGCTGGGGGCTGAGAGACAGACAGGAACCCACAATCAGGAGGCAACCCCTGGCCTGCAAGAGGAAGA
CAGAGGCTCCCAGGGCCGGTGCCCTGTGTGCCCCTGCACCAAGGCCGCTGAATAAGCCTGCCCTTACCCCCCTA
AGGGCTCCTTGCCCAATGCCAAGTGTGGGGATTCTGTGTCAGCAAGCCCTGTGGCTCCAGTGACGGTATTTCTAA
AGCCAACTTAGTTACCTAGAAATTAGCGCCATGTTGGAAACTGTGTCAGCAGCCCGGGCTGCACAGTGTGTAG
CCCAGCTCCAGGTCCACGGAGTGGTGTGGACCTCCACCTCACAGCTGCCTCTGGCAGCCAAGCCTCTTTTCGC
CCGGCCCCAGCCCCCTCTGGTTGATAAACGGGTGGGCCTCCTCAGCAGCGTGGCTGCCTTTCACCTTGATTTCCCC
AGGGCTCTCGGCAACATCGATAAACAGCCTCGCCACCAGCTGGGCCCTCCCCACCCAGTCTGCCAGGCTGGG
AGCTGGAGCTTGCTGAGTCTTGAATGCCCTTCTAGATGGCTTCTCTAGAGGCTCTCCTGGCAAGAGAGGGTCCCA
AGGGGAGCCCTGCAAAGCAAAGGCTCCTTGCTGCGGGCGGATAGAGAATCTCGCCTCTGTCTGGTGTGTTACCT
ACTGGGGGCACAGGAACAATTTCTCAAGGAGACAGTGGCATGGAGCTTTGAAAGACGAGTAGGTGTTAGCAAGG
AAATAAGGAGGAACGGGGGTACGGGCAGAGGAGAAAGCACATGCCAAGTCAGCAAAGAAAAGTAGAATTCGAAA
ACTTTTTAAAAATATTACTAAGGATTTTCACAATGCTGCACTGGGCTAGAACTGAAGCTAAACAGATACGTGG
TCCCTGCTGCTATGGGGCTTCCGTTCTAGAGGCAAGGACAGGTTGTGATGAGGGTTCTGAAGGATAGAGACCAAG
CAGGGAGGGTGTGAGGAGGCTTCTGCGAGACCTGAAGGATGGGAAGCCAGGAAGTGGGAGGGGTGGGGTCCAG
GCTGGAGGGGCCCAATGTAGGTGTAGAGGACTACAGCCCTGAGGGGCTGCTCCATGCGGCATTCTTGAGGTTCC
AAGAGGGGCAGCGCCACCTTGGGCCAGGCTCCTCTCCAGCAGCCTTGGTATGGGGTGGGGTGGGAAGACCCCTG
ATGCAGCCTGTCTCTGGGTGTGGGGGTGGAGGAGAGGGCTCCAGGGGACCCAGCCAGCCTTGACCCTGGAGGAA
AGTCAGGTGGGCTCAGAGAGAGCCCTCAAATCTGGGCCCTGGGTCAGGGTGGGGTCAAGTCCAGCCTGAAGAGA
ATGGACTCTGGAATCTGACGTCTGGGTTTGAATCCAGCTTCCAACACTCACTGGCTGTGTGACTTTGGGCAAGAT
AATTATCCCTTGTCCCCATGTGTGATTGGAGATAAACTACCCACCTACCAGATTGTTGTACCATGCTGGGCAA
ACACTAAGTGCTTAATAATGGTAGCCCACTGCTC

566/6881
FIGURE 526

MHRDQPHLECSQTRCPSAHTAPGHRLPIELSAPKPHGLFMHGDRPLDVQPCHPLPQSSLRAPASVPWAGEGEACL
HLPNSSSSRKQGLVCQSCRVGHSVGRSQPGLESPLIRPAGFYKDQSC

567/6881
FIGURE 527

AGCCGGATGGTCCCGCAGCTCGGGGCCGGCCATGCTTCGCGGTCCGTGGCGCCAGCTTTGGCTCTTTCTCCTGCT
GCTGCTCCCGGGCGCGCCTGAGCCCCGCGGCGCCTCCAGGCCGTGGGAGGGAACCGACGAGCCGGGCTCGGCCTG
GGCCTGGCCGGGCTTCCAGCGCCTGCAGGAGCAGCTCAGGGCGGCGGTGCCCTCTCCAAGCGGTACTGGACGCT
CTTCAGCTGCCAGGTGTGGCCCGACGACTGTGACGAGGACGAGGAGGCAGCCACGGGGCCCCCTGGGCTGGCGCCT
TCCTCTGTTGGGCCAGCGGTACCTGGACCTCCTGACCACGTGGTACTGCAGCTTCAAAGACTGCTGCCCTAGAGG
GGATTGCAGAATCTCCAACAACCTTTACAGGCTTAGAGTGGGACCTGAATGTGCGGCTGCATGGCCAGCATTGGT
CCAGCAGCTGGTCCTAAGAACAGTGAGGGGCTACTTAGAGACGCCCCAGCCAGAAAAGGCCCTTGCTCTGTCGTT
CCACGGCTGGTCTGGCACAGGCAAGAACCTTCGTGGCACGGATGCTGGTGGAGAACCCTGTATCGGGACGGGCTGAT
GAGTGAAGTGTGTCAGGATGTTTCATCGCCACGTTCCACTTTTCTCACCCTCAATATGTGGACCTGTACAAGGAGCA
GCTGATGAGCCAGATCCGGGAGACGCAGCAGCTCTGCCACCAGACCCTGTTTCATCTTCGATGAAGCGGAGAAGCT
GCACCCAGGGCTGCTGGAGGTCTTGGGCCACACTTAGAACGCCGGGCCCCCTGAGGGCCACAGGGCTGAGTCTCC
ATGGACTATCTTTCTGTTTTCTCAGTAATCTCAGGGGCGATATAATCAATGAGGTGGTCTAAAGTTGCTCAAGGC
TGGATGGTCCCGGAAGAAATTACGATGGAACACCTGGAGCCCCACCTCCAGGCGGAGATTGTGGAGACCATAGA
CAATGGCTTTGGCCACAGCCGTCTTGTGAAGGAAAACCTGATTGACTACTTCATCCCCTTCCTGCCTTTGGAGTA
CCGTACAGTGAGGCTGTGTGCACGGGATGCCTTCCTGAGCCAGGAGCTCCTGTATAAAGAAGAGACACTGGATGA
AATAGCCCAGATGATGGTGTATGTCCCAAGGAGGAACAACCTCTTTTCTTCCAGGGCTGCAAGTCTATTTCCCA
GAGGATTAACCTACTTCCGTGTCATGAAGGCTAGAGGAAGACTTCCTGGAAGTGCCTTTCTTCCACTAACAGGACCC
TGGGACCTGTAGGAGCACCCCGTTTGGGACTGTGAGGTGTTGAGGGTGTGGACTGGCATCCAGCAGCCACTAAC
AAACACACAACCTGGTGTGTAAGGAGGAGCCTTACATTAGAAGCCAAGCCAATCCTTTTTCTTTTTTTTGGAGGT
CCCACCGAGATAGATAGGAACCTGGATTGCTGAATTCAAAAACAGAGCCCATCTTAAGATCACTTGGTGCCTTA
AAGACACGCATTCCAAAGTGGAATGTGGTTGAAGAAAGTGGGCCAGGTGGTTGAAGAAAGCCATGTGGGAGCTCA
GCAAATCCCAAGGGCTTATTATGACACTCCAGATGGTCTCCTTAGCATCTCAGCTCTTCTGCAAGGAAGAGCTG
GGTGTAGGCCTCAGAGGCTGTAGGGTCCTTGGGTTACAGAGCCGGGAGAACGAAGTTCTGTGACCCAGGGGTG
GAGAATACACTCTAGGTTTGGGGCTGGTGGGCTTTCAAATTGGTACTTCCAGAGGAAGCCAAGCTGCTTCTGT
TGTGAGCGAATCAGCCAAGAGCCTGAGGCTGAAGGGAAGTACACAGAGGAAGATATTTTACAAACCAGGTCAG
TGTAGGCCAAGACTTATGGTCTACAGATTTTGGCGGGGAGGGGGACCTTTTCAAAGACAATAGGGGGTCTTGA
CATGTTTGTGTATGTAAAGATGATAAGATTAAATTTTGTATTTTCTAAAAA

568/6881

FIGURE 528

MLRGPWRQLWFLLLLLLPGAPEPRGASRPWEGTDEPGSAWAWPGFQRLQEQLRAGALS KRYWTLFSCQVWPDDC
DEDEEAATGPLGWRLPLL GQRYLDLLTTWYCSFKDCCPRGDCRISNNFTGLEWDLNVR LHGQHLVQQVLRTVRG
YLETPQPEKALALS FHGWSGTGKNFVARMLVENLYRDGLMSDCVRMF IATFHFPHPKYVDLYKEQLMSQIRETQQ
LCHQTLFIFDEAEKLHPGLLEVLGPHLERRAPEGHRAESPWTIFLFLSNLRGDI INEVVLKLLKAGWSREEITME
HLEPHLQAEIVETIDNGFGHSRLVKENLIDYFIFFLPLEYRHVRLCARD AFLSQELLYKEETLDEIAQMMVYVPK
EEQLFSSQGCKSISQRINYFLS

569/6881
FIGURE 529

ATTTCTCCATGTGGCAGACAGAGCAAAGCCACAACGCTTTCTCTGCTGGATTAAAGACGGCCACAGACCAGAAC
TTCCACTATACTACTTAAAATTACATAGGTGGCTTGTCAAATTCAATTGATTAGTATTGTAAAAGGAAAAAGAAG
TTCCTTCTTACAGCTTGGATTCAACGGTCCAAAACAAAAATGCAGCTGCCATTAAAGTCACAGATGAACAACTT
CTACACTGATTTTTAAAAATCAAGAATAAGGGCAGCAAGTTTCTGGATTCACTGAATCAACAGACACAAAAAGCTG
GCAATATAGCAACTATGAAGAGAAAAGCTACTAATAAAATTAACCCAACGCATAGAAGACTTTTTTTCTCTTCT
AAAAACAACCTAAGTAAAGACTTAAATTTAAACACATCATTTTACAACCTCATTTCAAAATGAAGACTTTTACCTG
GACCCTAGGTGTGCTATTCTTCTACTAGTGGACACTGGACATTGCAGAGGTGGACAATTCAAAATTAAAAAAT
AAACCAGAGAAGATACCCCTCGTGCCACAGATGGTAAAGAGGAAGCAAAGAAATGTGCATACACATTCTGGTACC
TGAACAAAGAATAACAGGGCCAATCTGTGTCAACACCAAGGGGCAAGATGCAAGTACCATTAAAGACATGATCAC
CAGGATGGACCTTGAAAACCTGAAGGATGTGCTCTCCAGGCAGAAAGCGGGAGATAGATGTTCTGCAACTGGTGGT
GGATGTAGATGGAAACATTGTGAATGAGGTAAAGCTGCTGAGAAAGGAAAGCCGTAACATGAACCTCTCGTGTAC
TCAACTCTATATGCAATTATTACATGAGATTATCCGTAAGAGGGATAATTCACTTGAACCTTCCCAACTGGAAAA
CAAAATCCTCAATGTCACCACAGAAATGTTGAAGATGGCAACAAGATACAGGGAAGTAAAGGTGAAATACGCTTC
CTGACTGATCTTGTCAATAACCAATCTGTGATGATCACTTGTGGAAGAACAGTGTGAGGATATTTCCCG
ACAAGACACCCATGTGTCTCCCCACTTGTCCAGGTGGTGCCACAACATATTCTAACAGCCAACAGTATACTCC
TGGTCTGCTGGGAGGTAACGAGATTCAGAGGGATCCAGGTTATCCAGAGATTTAATGCCACCACCTGATCTGGC
AACTTCTCCACCAAAAGCCCTTTCAAGATACCACCGGTAACCTTTCATCAATGAAGGACCATTCAAAGACTGTCA
GCAAGCAAAAGAAGCTGGGCATTTCGCTCAGTGGGATTATATGATTAAACCTGAAAACAGCAATGGACCAATGCA
GTTATGGTGTGAAAACAGTTTGGACCCTGGGGGTTGGACTGTTATTCAGAAAAGAACAGACGGCTCTGTCAACTT
CTTCAGAAATTGGGAAAATTATAAGAAAGGGTTTGGAAACATTGACGGAGAATACTGGCTTGGACTGGAAAAATAT
CTATATGCTTAGCAATCAAGATAATTACAAGTTATTGATTGAATTAGAAGACTGGAGTGATAAAAAAGTCTATGC
AGAATACAGCAGCTTTCGTCTGGAACCTGAAAGTGAATTCTATAGACTGCGCCTGGGAACCTTACCAGGGAAATGC
AGGGGATTCTATGATGTGGCATAATGGTAAACAATTCACCACACTGGACAGAGATAAAGATATGTATGCAGGAAA
CTGCGCCCACTTTCATAAAGGAGGCTGGTGGTACAATGCCTGTGCACATTCTAACCTAAATGGAGTATGGTACAG
AGGAGGCCATTACAGAAGCAAGCACCAAGATGGAATTTCTGGGCCGAATACAGAGGCGGGTCATACTCCTTAAG
AGCAGTTCAGATGATGATCAAGCCTATTGACTGAAGAGAGACACTCGCCAATTTAAATGACACAGAACCTTTGTAC
TTTTAGCTCTTAAAAATGTAAATGTTACATGTATATTACTTGGCACAAATTTATTTCTACACATAAAGTTTTTAA
AATGAATTTTACCGTAACCTATAAAAGGGAACCTATAAATGT

570/6881
FIGURE 530

MKTFTWTLGVLFLLVDTGHCRCGGQFKIKKINQRRYPRATDGKEEAKKCAYTFLVPEQRITGPICVNTKGQDAST
IKDMITRMDLENLKDVLRSQKREIDVLQLVVDVDGNIVNEVKLLRKESRNMNSRVTQLYMQLLHEIIRKRDNSLE
LSQLENKILNVTTEMLKMATRYRELEVKYASLTDLVNNQSVMITLLEEQLRIFSRQDTHVSPPLVQVVPQHIPP
SQQYTPGLLGGNEIQRDPGYPRDLMPPDLATSPTKSPFKIPPVTFINEGPFKDCQQAKEAGHSVSGIYMIKPEN
SNGPMQLWCENSLDPGGWTVIQKRTDGSVNFFRNWENYKKGFGNIDGEYWLGLENIYMLSNQDNYKLLIELEDWS
DKKVYAEYSSFRLEPESEFYRLRLGTYQGNAGDSMMWHNGKQFTTLDRDKDMYAGNCAHFHKGGWWYNACAHSNL
NGVWYRGGHYRSKHQDGIFWAEYRGGSYSRAVQMMIKPID

571/6881

FIGURE 531

ATGGGGCTGCTCACTGGCACTGGAGGCCGAAAGAAGGTGAGGGCAAGTGAGATGCAGGGACCTCTTCTCACAGCT
GTGCTGACAAGCACAAAAGGAGCCCCCACGTCCACACCAGGAAAGGGGGGATGGTGCACAAGATCCAGAGGCG
TCAACAGACACGTCCAGCGAACAAGCACAGGGTGGGCCGTCCGGGAGCCTTGCGCCAGGAGGAAAGCCGCAC
CGAGGGCAGCAGGGAGGCTCTGATGGCTGCCAAGAGGGTCTGATCCAGAAATATGGCCTCAATATGTGCCCCAG
TGTTTCCGTCAGTACGCGAAGGATATCGGTTTCATTAAATTGGACTAA

572/6881
FIGURE 532

TAGGTTTCATTTGGAAGCTTGAGTTTTATCATTGGCAACAAATGCCATCAGTTTTTTCCAAGCTGACTTGTTCTT
TTTTGAGCAAATCTCTGCCAAAATTACCCCTAAAATATGAATAATCAATTTGTCAGGTTTTTTCAAACATAAAAT
GTTCTGTGAAAACAAGCAGCTAGTTTCAGCTTGATCTCAAACAATTGTAACACTTTTTCTAGGGAAAAAAAAT
CTAGACATCCATACTTTATGGCAGATGTGCTTCATGCATACTTCCCATTTTGTACACAGAAATCAAAATAACG
TGTACTCAAGGTCAAGATTTACTAACAACTAAAATTCCAAAACCTGCTAAACAACAGAATGCTGAATAAGGTGATC
CAATTTTAAATATTTTATTTGAGACAACATCTCACCCTGTCAACCAGGCTGGAATCCAGTGGTGTGACACAGCTC
ACTGCATCCTCAACCTCCTGGGCTCAAGCGATCGTCCCACCTCAGTCTCATGAGTGGTTGGGACTACAGGCATGC
ACCATCATGCCCCGGCTCACTTTTTTTTTTCTTTTTTTTTTAAATTGTAGACACAGGGTCTCACTATGTTGCCAGG
CCGGTCTTGAACCTCTGGACTCAAGCAATCCCCCACCTTGGTTTCCCAAAGTGTGGGATTACAGGCATGAGCT
ACTACATACAACTGAAATCTTTTAAATATGCTTGTCAAAAAGTTTCTGATTACGCAAATGGCACTTATATATAAA
AGAGAGAACAAGAAGCTGGATGCGGTGGCTCACGCCTGTAATCCCAGCACTTTGGGAGGCTAGGTGGGCAGATCA
CGAGGTCAAGGATTCGAGACCAGCCTGGCCAACGTGGTGAACCCCCCTCTCTACTAAAAATACAAAAATAGCCAG
GCATGGTGGCCACGCCTGTAATCCCAGCTACTTGGAAAGGCTGAGGCAGGAGAATCGCTTGAACCTGGGAGGTGG
AGGTTGCAGTGAGCCAAGATTGCGCCGCTACATTCCAGCCTGGGCAACACGAGCGAAACTCCGTCTCAAAAAAAA
AAAAAAAAGAATAGAACTTGGCTGGATGCAGTGGCTCACACCTGTAATCCCAGCACTTTGGGAGGGCAGGGCAG
TTGGATCACTTGGGCCAGGAGTTCAAGACTAGCCTGGCTGACATGGTGAAACTCCATCTCTACTAAAAATACAG
AAAAAAAATTAGCTGGGAGTGGTGCCACACACCTGTCTATCCCAGCTACTCAGGTAGTTGAGGCACAACAATCAC
TTGAGCCTAGGAGGCCAGGCTGGAGTGAGGTGGCACGATCTTGGTTCACTGCAACCTCCACCTCCAGGGTTTACG
AGTAGCTGGGACTACAGGCCACACTACCACACCCGGCTAAATTTTCGCTTTTTTTCAGTAGAGACCGGGTTTTGC
CATGTTGGCCAGGCTGGTCTCGAAGCCTGAGCCATAACCTGGCGAAAAGATACTTTGAAATGTAACATCGTTTTT
CACTTCAAAATCAAGGTATTTTACCAACTTGCTTTTGGTACCTCCCCAAAACCTGATCTACTACATTTGCAGTTT
CATTCATTTTTCCAACATTAAGGAATAAACTTGAAGCGCTTACATTCTTGTGGGGGAAGTAAGAACAAAACACAT
AATAGGTCATGGTGGTCACATGAAAAGTAAAGGGTAACCGATTGGGGTAGGTGGTGGTAGTTTTGTGTGTGTGTT
TTTTTTTGAGACCAAGTCTCGCCGTTGCCAGGCTGGAGTGCGGTGGCATGATCTCAGTTCACTGCAACCTCCAC
TTCCTAGGTTCAAGCAATTCTCCTGCCTCAGCTACTCAAGAGGCTGAGGCAGGAGAATTCCTTGAACCTGGGAGG
CAGAGGTTGCAGTGAAGTATAGTGCCGCTGCCTCATTCTGGGCAAGAGCTAGACTCTTGTCTAAAAA
AAAAATACTTGAAGTCTATACCTTTGCTTCTTTAAACTTGCTCCACACAGTGTAGTCAAGCCGACTCTCCA
TACCTGTAGAAATTTAATCTCCTTTACCAATAGGTAGTAATAGGTGGATTTCTAAAAGATTTCCCAACTACTGTT
TCAGTTTAAAGATTAATCTCCATTTTCTTCTCAGAGAGATTCCCATCTGCTTTAATTTTAAATCATCACTAACA
GTAGCTCAATCAATTAACGTTAACATCAATAAACATGTACCTTTAAAAGGTATGACAGGAAGTGTCTTCATGTC
CTTACCCAAGCAAGTCATCCATGGATAAAAACGTTACCAGGAGCTGGAATACAAATGAGGACTTCCATTAAATG
CTAAACATCATTAACCTTTTTCAAAGTTGTCTGTC

573/6881

FIGURE 533

MCFMHTSHFVTQNIKITCTQGQDLLTTKIPKLLNNRMLNKVIQFLIFYLRQHLLTSPRLESSGVTQLTASSTSWA
QAIVPPQSHEWLGLQACTIMPGSLFFSFFF

574/6881

FIGURE 534

ATCTGTAGATGGGAGTGATTAGCTGTTTTAAAGTTAAATGTGACTGAGAAGGAAATTGAGTAGGGCAAATTTTA
AATGGGTATTATTTTCATCTTCAAACAGGCAGACCTGTTATCCTAAACTAGGTGAGTCAGCTTTTGGTACATGT
GATGATTTTCAGTGTAACCAATGATGTAATGATTCTGCCAAATGAAATATAATGATATCACTGTAAAACCGTTCC
ATTTTGATTCTGAGGTTACTCTACTAACAAGCATCACACATTTGTATTTTGCCCTGATTAATATGTTGGCTTCGC
TTTCAGGGTTTTTAATGACCACAACAAGCAAGCATGCAGCTTACTGCTTGAAAGGGTCTTGCCTCACCCAAGCTA
GAGTGCAGTGGCCTTTGAAGCTTACTACAGCCTCAAACCTCTGGGCTCAAGTGATCCTCAGCCTCCCAGTGGTCT
TTGTAGACTGCCTGATGGAGTCTCATGGCACAAGAAGATTAAACAGTGTCTCCAATTTTAATAAATTTTGCAA
TCC

575/6881
FIGURE 535

AGTTAAACAGCTGAGCTTCTGAATGCCTGCAAGAAGCTGCCCTTTGAAATTAAGAACTTCGTGAAGAAAACAGA
GGCTCTTCGGTTGCAGTATCGCTACTTAGACTTGCGTAGTTTCCAAATGCAGTATAACCTGCGACTGAGGTCCCA
GATGGTCATGAAAATGCGGGAATATCTCTGTAATCTGCATGGGTTTGTGGATATAGAAACCCACATTGTTTAA
GAGGACCCAGGGGTGCCAAAGAGTTTTTAGTACCATCCAGGGAACCTGGAAAGTTTTGTTCTCTCCCTCAGAG
TCCTCAACAGTTTAAAGCAACTTCTGATGGTTGGCGGTTTAGACAGATATTTTCAGGTTGCCGATGTTATCGAGA
TGAAGGTTCAAGACCAGACAGACAGCCTGAGTTTACTCAGATTGACATAGAGATGTCATTTGTAGACCAGACTGG
GATCCAGAGTTTAAATTGAGGGTTTGCTCCAGTATTCCTGGCCCAATGACAAAGATCCTGTGGTTGTTCTTTTCC
TACTATGACTTTTGCTGAGGTGCTGGCCACCTATGGAACTGATAAACCTGACACTCGCTTTGGAATGAAGATTAT
AGATATCAGTGATGTGTTTAGAAACACAGAGATTGGATTCTTCAAGATGCACTTAGTAAGCCCCATGGAAGTGT
GAAAGCCATATGTATCCCTGAAGGAGCAAAATACTTAAAAAGGAAAGACATTGAATCCATTAGAACTTTGCAGC
TGACATTTTAAATCAGGAAATCTTACCTGTATTCTTAACGCCAATAGAACTGGAATTCTCCAGTTGCTAATTT
CATAATGGAGTCACAAAGACTGGAATTAATCAGACTAATGGAGACCCAAGAGGAAGATGTGGTCTCTAACTGC
TGGAGAGCACAAATAAGCATGCTCTTTGTTAGGAAATACGACTGGAATGTGCTGACCTCTAGAAACAAGAGG
AGTGGTGCTCCGTGACCCCACTCTGTCTCTTTCTTTGGGTGGTAGATTTCCCACTCTCTCCTGCCCAAGGAGGA
AAGTCCCAGAGAGCTGGAATCGGCCCACCACCCATTTACTGCTCCCCACCCAGTGACATACATCTCCTGTACAC
TGAGCCCAAAAGGCCCGTAGCCAACACTATGACTTGGTTTTAAATGGCAATGAAATAGGAGGTGTTCAATTCCG
AATTCACAATGCAGAGCTGCAGCGTTATATCCTGGCAACCTTACTAAAGGAGGATGTGAAAATGCTCTCCCATCT
GCTCCAGGCTTTAGATTATGGGGCACCCCTCATGGAGGAATTGCCCTTAGGGTTAGACAGACTGATATGCCTTGT
CACTGGATCTCCAAGCATCAGAGATGTCATAGCCTTCCCAAAGTCTTCCGGGGACATGACCTCATGAGCAATAC
CCCAGATTCTGTCCCTCCTGAGGAACTGAAGCCCTATCATATCCGAGTCTCCAAGCCAACAGACTCCAAGCAGA
AAGAGCTCATTTGAATCATGCATACCATGCAGAAAGTTGAGCTTTTAGGTTTTGTCCTCTTTGCTTCCCCAAGGCT
AAAGTCAGATCTAGAGTTCTGCCACAGGTCTAACAATCAAGTCTTTAGATGGAAGGAATCCAGGCAACATTCTTC
ACCACAACGAAGAAACAGATAAAAGATACCCAATTTTGACTTGATTTCATGCATCATTGGAATTTTTTTGGTTA
GGACTTTTTTTGAAGTTCCTTTTTTACTTAGGTGTGAAAGATGGTTCTTTGTTGAAATAATATAGTGGTTTAGTGT
TTTCAAATCATGTTTCTCATACCCAGATAGTAGATTATTCACTTAGGACAGAGGTAATCAAATTATGTGTGAAAT
GTAGGAAAATGCTTGCCCTGTAACTAGTGAGTTGATGGAGCATTGCTTCATCATCCTCATCAAGAGAATCAT
ATAAATTAAGCTTTATAATGACATTCAACCATCAACATAATATAGTGAGGAGTAGCATAATATTTTTTAATAAT
GCAGAAAACATCACTGAAATGAGAGTCACAAATTTTTCTCAGTGTTTCAGCCTGAGTAAGTTACATAAACCTCG
CTTAGCCTCCCTTCCTGCTAATGTGTAAATACATACTTGCCCTGGCTACCTCACCGGGCTGTTATTGCTGGAAT
CAGAGGAGATAACATATATGGAAGATAAAGTGAATAAAAGTACTTTGAAAACT

576/6881

FIGURE 536

MQYNLRRLRSQMVMKMREYLCNLHGFVDIETPTLFKRTPGGAKEFLVPSREPGKFCSLPQSPQQFKQLLMVGGLDR
YFQVARCVRDEGSRPDRQPEFTQIDIEMSFVDQTGIQSLIEGLLQYSWPNDKDPVVVPFPTMTFAEVLATYGTDK
PDTRFGMKIIDISDVFRNTEIGFLQDALSKEPHGTVKAICIEGAKYLKRKDIESIRNFAADHFNQEILPVFLNAN
RNWNSPVANFIMESQRLELIRLMETQEEDVLLTAGHEHNKACSLGKLRLECADLLETGRGVVLRDPTLFSFLWVV
DFPLFLPKESPRELESAAHHPFTAPHPSDIHLLYTEPKKARSQHYDLVLNGNEIGGGSIRIHNAELQRYILATLL
KEDVKMLSHLLQALDYGAPPHGGIALGLDRLICLVGTGSPSIRDVIAFPKSFRGHDLMSTPDSPPEELKPYHIR
VSKPTDSKAERAH

577/6881
FIGURE 537

GCCTGTCTGCATTCTACTATATAAAGCAGCAGAGACGTTGACTAGCGCATATTTGCTAAGAGCACCATGCGCGCA
GCAGCCATCTCCACTCCAAAGTTAGACAAAATGCCAGGAATGTTCTTCTCTGCTAACCCAAAGGAATTGAAAGGA
ACCACTCATTCACTTCTAGACGACAAAATGCAAAAAGGAGGCCAAAGACTTTTGGAAATGGATATGAAAGCATAC
CTGAGATCTATGATCCCACATCTGGAATCTGGAATGAAATCTTCCAAGTCCAAGGATGTACTTTCTGCTGCTGAA
GTAATGCAATGGTCTCAATCTCTGGAAAACTTCTTGCCAACCAAACCTGGTCAAAATGTCTTTGGAAGTTTCCTA
AAGTCTGAATTCAGTGAGGAGAATATTGAGTTCTGGCTGGCTTGTGAAGACTATAAGAAAACAGAGTCTGATCTT
TTGCCCTGTAAAGCAGAAGAGATATATAAAGCATTTGTGCATTGAGATGCTGCTAAACAAATCAATATTGACTTC
CGCACTCGAGAATCTACAGCCAAGAAGATTAAAGCACCAACCCACGTTTGTGATGAAGCACAAAAGTCATA
TATACTCTTATGGAAGGACTCTTATCCCAGGTTCTCTCAAATCAGATATTTACTTAAATCTTCTAAATGACCTG
CAGGCTAATAGCCTAAAGTGACTGGTCCCTGGCTGAAGGGAATTAACAGATAGTATCAAGCGCAGAAGGAATGTG
CCAGTATGGCTCCCTGGGTGAACAGCTTGGCCTTTTTTGGGTGCTTGTGACAGGCCAAGAAGAACAAATGACTCAG
AATGGATTAAACATGAAAGTTATCCAGGCGCAGAGTTGAAGAAGCATAAGCAAGACAAAACAGAGAGACCGCAGA
AGGAGGAAGATACTGTGGTACTGTCATAAAAAACAGTGGAGCTCTGTATTAGAAAGCCCCCTCAGAACTGGGAAGG
CCAGGTAACCTCTAGTTACACAGAACTGTGACTAAAGTCTATGAACTGATTACAACAGACTGTAAGAATCAAAG
TCAACTGACATCTATGCTACATATTATTATATAGTTTGTACTGAGCTATTGAAGTCCCATTAACCTAAAGTATAT
GTTTTCAAATTGCCATTGCTACTATTGCTTGTGCGGTGTTATTTTATTTTATTGTTTTGACTTTGGAAGAGATGA
ACTGTGTATTTAACTTAAGCTATTGCTCTTAAACCAGGGAGTCAGAATATATTTGTAAGTTAAATCATTGGTGC
TAATAATAAATGTGGATTTTGTATTAAATATATAGAAGCAATTTCTGTTTACATGTCCTTGCTACTTTTAAAAA
CTTGCATTTATTCCCTCAGATTTTAAAAATAAATAAATAATTCATTT

578/6881
FIGURE 538

CTTTTCGATCCGCCATCTGCAGTGGAGCCGCCACCAAAATGCAGATTTTCGTGAAAACCCTTATGGGGAAGACCA
TCACCCTCGAGGTTGAACTCTCGGATACAATAGATAATGTAAAGGCCAAGATCCAGGATAAGGAAGGAATTCCTC
CTGATCAGCAGAGACTGATCTTTGCTGGCAAGCAGTTGGAAGATGGACGTACTTTGTCTGACTACAATATTCAA
AGGAGTCTACTCTTCATCTTGTGTTGAGACTTCGTGGTGGTGCTAAGAAAAGGAAGAAGAAGTCTTACACCACTC
CCAGGAAGAATAAGCACAAGAGAAAGAAGGTTAAGCTGGCTCTCCTGAAATATTATAAGGTGGATGAGAATGGCT
TTATGGCAAGCCACTTTGACAGACATTATTGTGGCAAATGTTGTCTGACTTACTGTTTCAACAAACCAGAAGACA
AGTAACTGTATGACTTAATAA

579/6881
FIGURE 539

MQIFVKTLMGKTIITLEVELSDTIDNVKAKIQDKEGIPPDQORLIFAGKQLEDGRTLSDYNIQKESTLHLVLRRLRG
GAKKRKKKSYYTPRKNKHKRKKVKLALLKYYKVDENGFMASHFDRHYCGKCCLTYCFNKPEDK

580/6881
FIGURE 540

AAAACAGCCGGGGCTCCAGCGGGAGAACCATAATGCAAAGTGCTATGTTCTTGGCTGTTCAACACGACTGCAGAC
CCATGGACAAGAGCGCAGGCAGTGGCCACAAGAGCGAGGAGAAGCGAGAAAAGATGAAACGGACCCTTTTAAAAG
ATTGGAAGACCCGTTTGGAGCTACTTCTTACAAAATTCCTCTACTCCTGGGAAGCCCCAAAACCGGCAAAAAAAGCA
AACAGCAAGCTTTTCATCAAGCCTTCTCCTGAGGAAGCACAGCTGTGGTCAGAAGCATTGACGAGCTGCTAGCCA
GCAAATATGGTCTTGGCTGCATTTCAGGGCTTTTTTAAAGTCGGAATTCTGTGAAGAAAATATTGAATTCTGGCTGG
CCTGTGAAGACTTCAAAAAAACCAAATCACCCCAAAAGCTGTCCTCAAAGCAAGGAAAATATATACTGACTTCA
TAGAAAAGGAAGCTCCAAAAGAGATAAACATAGATTTTCAAACCAAACTCTGATTGCCCAGAATATACAAGAAG
CTACAAGTGGCTGCTTTTACAACGCCCAGAAAAGGGTATACAGCTTGATGGAGAACAACCTTTATCCTCGTTTCT
TGGAGTCAGAATTCTACCAGGACTTGTGTAAAAAGCCACAAATCACCACAGAGCCTCATGCTACATGAAATGTAA
AAGGGAGCCCAGAAATGGAGGACATTTTCTTTTCTGAGGGGAAGGACTGTGACCTGCCATAAAGACTGAC
CTTGAATTCAGCCTGGGTGTTTTCAGGAAACATCACTCAGAACTATTGATTCAAAGTTGGGTAGTGAATCAGGAAGC
CAGTAACTGACTAGGAGAAGCTGGTATCAGAACAGCTTCCCTCACTGTGTACAGAACGCAAGAAGGGAATAGGTG
GTCTGAACGTGGTGTCTCACTCTGAAAAGCAGGAATGTAAGATGATGAAAGAGACAATGTAATACTGTTGGTCCA
AAAGCATTAAAAATCAATAGATCTGGGATTATGTGGCCTTAGGTAGCTGGTTGTACATCTTCCCTAAATCGATC
CATGTTACCACATAGTAGTTTTAGTTTAGGATTTCAGTAACAGTGAAGTGTTTACTATGTGCAAGGGTATTGAAGT
TCTTATGACCACAGATCATCAGTACTGTTGTCTCATGTAATGCTAAACTGAAATGGTCCGTGTTTGCAATTGTTA
AAAATGATGTGTGAAATAGAATGAGTGCTATGGTGTGAAAACGTCAGTGTCGGTTATGAGTGCCAAAAATCTGT
CTTGAAGGCAGCTACACTTTGAAGTGGTCTTTGAATACTTTTAATAAATTTATTTTGATAAATAATATTG

581/6881
FIGURE 541

MQSAMFLAVQHDCRPMDSAGSGHKSEEKREKMKRTLLKDWKTRLSYFLQNSSTPGKPKTGKKSQQAFIKPSPE
EAQLWSEAFDELLASKYGLAAFRFLKSEFCEENIEFWLACEDFKKTKSPQKLSSKARKIYTDFIEKEAPKEINI
DFQTKTLIAQNIQEATSGCFTTAQKRVSMLMENNSYPRFLESEFYQDLCKKPQITTEPHAT

582/6881
FIGURE 542

TGAAGATGAAAAAGAAGATCATAAAATGTGCGCCAACAACGGCAGGCGGCATCTAAAGCAGCTTCTAAACAGAG
AGAGATGCTCATGGAAGATGTGGGCAGTGAGGAAGAACAAGAAGAGGAGGATGAGGCACCATTCCAGGAGAAAGA
TTCCGGCAGCGATGAAGATTTCTAATGGAAGATGATGACGATAGTGACTATGGCAGTTCGAAAAAGAAAAACAA
AAAGATGGTTAAGAAGTCCAAACCTGAAAGAAAAGAAAAGAAAATGCCCAAACCCAGACTAAAGGCTACAGTGAC
GCCAAGTCCAGTGAAAGGCAAAGGGAAAGTGGGTGCCCCACAGCTTCAAAGGCATCAAAGGAAAAGACTCCTTC
TCCCAAAGAAGAAGATGAGGAACCGGAAAGCCCGCCAGAAAAGAAAACATCTACAAGCCCCCACCAGAGAAATC
TGGGGATGAAGGGTCTGAAGATGAAGCCCCTTCTGGGGAGGATTAAAAGTGATGATGGTCTGGGGAGAGATTTTA
TTAAAAAAGAAAAGAGGGAGGAAAAAAGAACCTACTTAAGATAGAACATGGTTTTGGCTATGGCTT
GACTCATGGGCTTTCAGTGCTTTTTTCCATTTGTTGAAAGTAACATTTCTCTCTCTCTCTTT

583/6881
FIGURE 543

AGCCTGAGGAGCTATTTTGAGCAATGGGGAATGCTCACGGACTGTGTGGTAATGAGAGACCCAAACACCAAGTGC
TCCAGGGGCTTTGGGTTTGTACATATGCCATTGTGGAGGAGGTGGATGCAGCCATGAATGCAAGGCCACACAAG
GTGGATGGAAGAGTTGGAGAACCAAAGAGAGCTGTTTCAAGGGAAGATTCTCAAAGACCAGGTGCTCACTTAACT
GTGAAAAAGATATTTGTTGGCAGCATTAAAGAAGACACTGAAGAACATCACCTAAGAGATAATTTGAACAGTTTG
GAAAAACGGAAGTGATTGAAATCATGACTGACTGAGGCAGTGGCAAGAAAAGGGGCTTTGCCTTTGTAACCTTTG
ATGACCATGACTCTGTGAATAAGACTGCCATTGAGAAATACCATCCTACGAATGGCCATAACTGTGAAGTTAGGA
AAGCCCTGTCAAAGTAAGAGATGGCTAGTGCTTCATCCAGCCAAAGAAGTCGAAGTGGTTCTGGAACTTTGGTT
TGGTCGTGGAGGTGGTTTCGGTTGGCATGACAGCCGTGGTGGTGGTGGTGGATATGGTGGCAGTGAGGATGGCGA
TAATGGATTTGGTAATGATGGAAGCAATTTTGGAGGTGGTGGGAAGCTACAGTGATTTTGGCAACTACAATAATCA
GTCTTCAAATTTTGGACCCATGAAGGGAGGAACTTTGGAGGCAGAAGCTCTGGGCCCTATGGCGGTGGAGGCC
AATACTCTGCCAAACCACGAAACCAAGGTGGCTATGGTGGTTTCAGTAGCAGCAGTAGCTATGGCAGTGGCAGAA
GATTTTAATTAGGAAACAAAGCTTAGCTGGAGAGGAGAGCCGGAGAAGTGACAGGGAAGCTACAGGTTACAACAG
ATTTGTGAAGTCAAGCAAGCAGTGGTGGCAGGGCCTAACTGCTACAAAGGAGACATGTTTATAGACAAATACTC
ATGTGTATGGGCAAAAACTCGAGGACTGTATTTGTGACTAATTGTGTAACAGGTTATTTAGTTTCTGTCTGT
GGAAAGCGTAAAGCATTCCAACAAAGGGTTTTAATGCAGATTTTTTTTTTTTGGCACCCATGCTGTTGATTGCT
AAATGTAATAGTCTGATGGTGTGCTGAATAAATGTCTT

584/6881
FIGURE 544

AGGCAGAGCCCGCGAGGAGGTGACGCGGCTGCGGAGGTGACGCGGGAGGTCGCGCGCCCCTTCCGGCGCGGGGAG
GGCGCTGAAGATCGGGGGCCGCTCGGCCGCGAGGCCGCTCCAGCGCCGCGGGATGTAGCGCGGGGACCGCGGGCCC
CCAGCAGAGCCCGCCTGCCCGGCTTTTAGAAGACCAAACTGGACAATGGACTTTGCTCACTATGATGACATGATC
TCCATTGACCTCCATTTGTCTACCATCAGAGGGAGATCTCTGCCCCCTGGGGCTGAGAGACCCCAACCTTTCCCC
AAGCTGAAGCTGCAGGGTATTGAGGTACCAGCCAGATGTCTTCCCAAAAGGATCTGTGGTGGCACAGGGGAATG
GGGCTCCTGCCAGTAACAGGGAAGCTGACACGGTGAAGTGGCTGAACTGGGACCCCTGCTAGAAGAGAAGGGCA
AACGGGTAAATCGCCAACCCACCCAAAGCTGAAGAAGAGCAAACATGCCCAGTGCCCCAGGAAGAAGAGGAGGAGG
TGCGGGTACTGACACTTCCCTGCAAGCCCACCACGCCATGGAGAAGATGGAAGAGTTTGTGTACAAGGTCTGGG
AGGGACGTTGGAGGGTCAATCCCATATGATGTGCTCCCTGACTGGCTAAAGGACAACGACTATCTGTACATGGTC
ATAGACCTCCCATGCCCTCCTTTGCGGCTTGCTTCAAGAGCATCTTCCGCATTTCATACAGAACTGGCAACATCT
GGACCCATCTGCTTGGTTTTCTGTCTGTTTCTCTTTTGGGAATCTTGACCATGCTCAGACCAAATATGTACTTCA
TGGCCCCCTCTACAGGAGAAGGTGGTTTTTGGGATGTTCTTTTGGGTGCAGTGCTCTGCCTCAGCTTCTCCTGGC
TCTTTCACACCGTCTATTGTCAATTCAGAGAAAGTCTCTCGGACTTTTTCCAACTGGACTATTACAGGATTGCTC
TTCTAATTATGGGGAGCTTTGTCCCTGGCTCTATTATTCTTCTACTGCTCCCCACAGCCACGGCTCATCTACC
TCTCCATCGTCTGTGCTCCTGGGCATTTCTGCCATCATTGTGGCGCAGTGGGACCGGTTTGCCACTCCTAAGCACC
GGCAGACAAGAGCAGGCGTGTCTCTGGGACTTGGCTTGAGTGGCGTCGTGCCCACCATGCACTTTACTATCGCTG
AGGGCTTTGTCAAGGCCACCACAGTGGGCCAGATGGGCTGGTTCTTCTCATGGCTGTGATGTACATCACTGGAG
CTGGCCTTTATGCTGCTCGAATTCCTGAGCGCTTCTTCTGGAAAATTTGACATATGGTTCAGTCTCATCAGA
TTTTCCATGTCTGGTGGTGGCAGCAGCCTTTGTCCACTTCTATGGAGTCTCCAACCTTCAGGAATTCGGTTACG
GCCTAGAAGGCGGCTGTACTGATGACACCCTTCTCTGAGCCTTCCCACCTGCGGGGTGGAGGAGGAACCTCCCAA
GTGCTTTTAAAAATAACTTCTTTGCTGAAGTGAGAGGAAGAGTCTGAGTTGTCTGTTTCTAGAAGAAACCTCTTA
GAGAATTCAGTACCAACCAAGCTTCAGCCCACTTTACACCCACTGGGCAATAAACTTTCCATTTCCATTCTCCT
AGCTGGGGATGGGGCATGGTCAAACCTTAGCCATCCCTCCTCAGCAAGGCATCTACCGGCCCTCACAGAGACAG
TACTTTGAACTCATGTTGAGATTTTACCCTCTCCTCCAACCATTTTGGGAAAATTATGGACTGGGACTCTTCAG
AAATTCTGTCTTTTCTTCTGGAAGAAAATGTCCCTCCCTTACCCCCATCCTTAACTTTGTATCCTGGCTTATAAC
AGGCCATCCATTTTGTAGCACACTTTTCAAAAACAATTATATACCCTGGTCCCATCTTTCTAGGGCCTGGATCT
GCTTATAGAGCAGGAAGAATAAAGCCACCAACTTTTACCTAGCCCGGCTAATCATGGAAGTGTGTCCAGGCTTCA
AGTAACTTGAGTTTTAATTTTTTTTTTCTTGGCAGAGTAATGTAAAATTTAATGGGGAAAGATATTTAATAT
TTAATACTAAGCTTTAAAAAGAAACCTGCTATCATTGCTATGTATCTTGATGCAAAGACTATGATGTTAATAAAA
GAAAGTACAGAAGACACTTGGCATTCAAAGATTTT

FIGURE 545

[illegible]

586/6881
FIGURE 546

MARGPGPLGRPRPDTVAMPKRGKRLKFRAHDACSGRVTVADYANS DPAVVRSGRVKKAVANAVQQEVKSLCGLEA
SQVPAAEEALSGAGEPCDIIDSSDEMDAQEESIHERTVSRKKKSKRHKEELDGAGGEEYPMDIWLLASYIRPEDI
VNFSLICKNAWTVTCTAAFWTRL YRRHYTL DASLPLRLRPESMEKLRCLRACVIRSLYHMYEPPAARISK NPAIP
ESTPSTLKNSKCLLFWCRKIVGNRQEP MWEFNFKFKKQSPRLKSKCTGGLQPPVQYEDVHTNPDQDCCLLQVTTL
NFI FIP IVMGMIFTLFTIN VSTDMRHHRVRLVFQDSPVHGGRKLRSEQGVQVILDPVHSVRLFDWWHPQYPFSLR
A

587/6881

FIGURE 547

AGTGGAGTGGGACAGGTATATAAAGGAAGTACAGGGCCTGGGGAAGAGGCCCTGTCTAGGTAGCTGGCACCAGGA
GCCGTGGGCAAGGGAAGAGGCCACACCCCTGCCCTGCTCTGCTGCAGCCAGAATGGGTGTGAAGGCGTCTCAAACA
GGCTTTGTGGTCCCTGGTGTCTGCCAGTGTCTCTGCATACAACTGGTCTGCTACTACACCAGCTGGTCCCAG
TACCGGGAAGGCGATGGGAGCTGCTTCCCAGATGCCCTTGACCGCTTCCTCTGTACCCACATCATCTACAGCTTT
GCCAATATAAGCAACGATCACATCGACACCTGGGAGTGAATGATGTGACGCTCTACGGCATGCTCAACACACTC
AAGAACAGGAACCCCAACCTGAAGACTCTCTTGTCTGTGCGAGGATGGAACCTTGGGTCTCAAAGATTTTCCAAG
ATAGCCTCCAACACCCAGAGTCGCCGGACTTTCATCAAGTCAGTACCGCCATTTCTGCGCACCCATGGCTTTGAT
GGGCTGGACCTTGCCCTGGCTCTACCTGGACGGAGAGACAAACAGCATTTTACCACCCTAATCAAGGAAATGAAG
GCCGAATTTATAAAGGAAGCCAGCCAGGGAAGAGCAGCTCCTGCTCAGCGCAGCACTGTCTGCGGGGAAGGTC
ACCATTGACAGCAGCTATGACATTGCCAAGATATCCCAACACCTGGATTTTATTAGCATCATGACCTACGATTTT
CATGGAGCCTGGCGTGGGACCACAGGCCATCACAGTCCCCTGTTCCGAGGTGAGGAGGATGCAAGTCCCTGACAGA
TTCAGCAACACTGACTATGCTGTGGGGTACATGTTGAGGCTGGGGGCTCCTGCCAGTAAGCTGGTGATGGGCATC
CCCACCTTCGGGAGGAGCTTCACTCTGGCTTCTTCTGAGACTGGTGTGGAGCCCCAATCTCAGGACCGGGAATT
CCAGGCCGGTTACCAAGGAGGCAGGGACCCCTTGCCCTACTATGAGATCTGTGACTTCCTCCGCGGAGCCACAGTC
CATAGAATCCTCGGCCAGCAGGTCCCCTATGCCACCAAGGGCAACCAGTGGGTAGGATACGACGACCAGGAAAGC
GTCAAAAGCAAGGTGCAGTACCTGAAGGACAGGCAGCTGGCGGGCGCCATGGTATGGGCCCTGGACCTGGATGAC
TTCCAGGGCTCCTTCTGTGGCCAGGATCTGCGCTTCCCTCTACCAATGCCATCAAGGATGCACTCGCTGCAACG
TAGCCCTCTGTCTGCACACAGCACGGGGGCCAAGGATGCCCGTCCCCCTCTGGCTCCAGCTGGCCGGGAGCCT
GATCACCTGCCCTGTCTGAGTCCCAGGCTGAGCCTCAGTCTCCCTCCCTTGGGGCCTATGCAGAGGTCCACAACAC
ACAGATTTGAGCTCAGCCCTGGTGGGCAGAGAGGTAGGGATGGGGCTGTGGGGATAGTGAGGCATCGCAATGTAA
GACTCGGGATTAGTACACACTTGTGATTAATGGAAATGTTTACAGATCCCCAAGCCTGGCAAGGGAATTTCTTC
AACTCCCTGCCCCCAGCCCTCCTTATCAAAGGACACCATTTTGGCAAGCTCTATACCAAGGAGCCAAACATCC
TACAAGACACAGTGACCATACTAATTATACCCCTGCAAAGCCCAGCTTGAAACCTTCACTTAGGAACGTAATCG
TGTCCCCTATCCTACTTCCCCTTCCTAATTCCACAGCTGCTCAATAAAGTACAAGAGCTTAACAGTGTGTTGGCG
CTTTGCTTTGGTCTATCTTTGAGCGCCCACTAGACCCACTGGACTCACCTCCCCATCTCTTCTGGGTTCCTTCC
TCTGAGCCTTGGGACCCCTGAGCTTGACAGATGAGGCCGCCATGT

588/6881
FIGURE 548

CAGGGTAAACGCTGTCTTGTGGACCCGCACTTCCCACCCGAGACCTCTCACTGAGCCCCGAGCCGCGCGACATGA
GCCACGGGAAGGGAACCGACATGCTCCCGGAGATCGCCGCCGCCGTGGGCTTCCTCTCCAGCCTCCTGAGGACCC
GGGGCTGCGTGAGCGAGCAGAGGCTTAAGGTCTTCAGCGGGCGCTCCAGGAGGCACTCACAGAGCACTACAAAC
ACCACTGGTTTTCCGAAAAGCCGTCCAAGGGCTCCGGCTACCGCTGCATTTCGCATCAACCACAAGATGGACCCCA
TCATCAGCAGGGTGGCCAGCCAGATCGGACTCAGCCAGCCCCAGCTGCACCAGCTGCTGCCAGCGAGCTGACCC
TGTGGGTGGACCCCTATGAGGTGTCTACCGCATTTGGGAGGACGGCTCCATCTGCGTCTTGTACGAGGAGGCC
CACTGGCCGCCTCCTGTGGGCTCCTCACCTGCAAGAACCAAGTGTGCTGGGCCGAGCAGCCCCCTCCAAGAACT
ACGTGATGGCAGTCTCCAGCTTAGGCCCTTCCGCCCCGCCCTGGGCGCCGCCGTGCTCATGCTGCCGTGACAACA
GGCCACCACATACCTCAACCTGGGGAAGTGTATTTTTAAATGAAGAGCTATTTATATATATTATTTTTTTTAAG
AAAGGAGGAAAAGAAACCAAAAGTTTTTTTTTAAGAAAAAAATCCTTCAAGGAGCTGCTTGGAAAGTGGCCTCCC
CAGGTGCCTTTGGAGAGAACTGTTGCGTGCTTGAGTCTGTGAGCCAGTGTCTGCCTATAGGAGGGGAGCTGTAA
GGGGGTAGACCTAGCCAAGGAGAAGTGGGAGACGTTTGGCTAGCACCCAGGAAGATGTGAGAGGGAGCAAGCAA
GGTTAGCAACTGTGAACAGAGAGGTGGGATTGCCCCGGGGGAGGAAGAGAGGCCAAGTTCAGAGCTCTCTGTC
TCCCCAGCCAGACACCTGCATCCCTGGCTCCTCTATTACTCAGGGGCATTATGCCTGGACTTAAACAATACTA
TGTTATCTTTTCTTTTATTTTCTAATGAGGTCTGGGCAGAGAGTGAAAAGGCCCTCTCCTGATTCTACTGTCC
TAAGCTGCTTTTCTTGAAATCATGACTTGTTTCTAATTCTACCTCAGGGGCCCTGTAGATGTTGCTTTCCAGCCA
GGAATCTAAAGCTTTGGGTTTTCTGAGGGGGGAGGAGGGAAGTGGAGGTTATTGGGGTTAGGATGGGAAAGGAA
TCTGCACAAAACCTTTGCTTTGCTAGTGTGCTTTGTGTGTATGTGTGGCAAATAATTTGGGGGTGATTGCAAT
GAAATTTTGGGACCCAAAGAGTATCCACTGGGGATGTTTTTTGGCCAAAACCTTCTCTTTTGGAAACCATGAAA
GTCTTGATGCTGCTGCCATGATCCCTTTGAGAGGTGGCTCAAAAGCTACAGGGAAGTCCAGGTCTTTTACTG
CCTTCTTTTCAAAAGCACAACTCTCCTCTAACCTCCCCCTCCCCCTTCCCTTCTGGTGGGTCATAGAGCTACCG
TATTTCTAGGACAAGAGTTCTCAGTCACTGTGCAATATGCCCTGGGTCCCAGGAGGGTCTGGAGGAAAAGT
GCTATCAGAACCTCCTGATGCCCTGGTGGGCTTAGGGAACCATCTCTCCTGCTCTCCTTGGGATGATGGCTGGCT
AGTCAGCCTTGCATGTATTCTTGGCTGAATGGGAGAGTGCCCCATGTTCTGCAAGACTACTTGGTATTCTTGTA
GGGCCGACACTAAATAAAAGCCAAACCTTGGGCACTGTTTTTCTCCCTGGTGCTCAGAGCACCTGTGGGAAAGG
TTGCTGTCTGTCTCAGTACAATCCAAATTTGTCGTAGACTTGTGCAATATATACTGTTGTGGGTTGGAGAAAAGT
GGAAAGCTACACTGGGAAGAACTCCCTTCCCTTCAATTTCTCAGTGACATTGATGAGGGGTCTCATAAGACCTC
GAGTTTCCCAAACCGAATCACCTTAAGAAGGACAGGGCTAGGGCATTGGCCAGGATGGCCACCCTCCTGCTGTT
GCCCTTAGTGAGGAATCTTACCCCACTTCCCTTACCCCCAGGTTCTCCTCCCCACAGCCAGTCCCCCTTCCCTG
GATTTCTAAACTGCTCAATTTTGACTCAAAGGTGCTATTTACCAAACACTCTCCCTACCCATTCTGCCAGCTCT
GCCTCCTTTTCAACTCTCCACATTTTGATTGCTTCCCAGACCTGCTTCCAGTCTTTATTGCTTTAAAGTTCA
TTTGGGCCACAGACCCAAAGAGCTAATTTTCTGTTTGTGGGTTGAAACAAAGCTGTGAATCACTGCAGGCTGTG
TTCTTGATCTTGTCTGCAACAGGTCCCTGCCTTTTTAGAAGCAGCCTCATGGTCTCATGCTTAATCTTGCTC
TCTTCTCTTTTATGATGTTCACTTTAAAAACAACAAACCCCTGAGCTGGACTGTTGAGCAGGCCTGTCTCTC
CTATTAAGTAAAAATAAATAGTAGTAGTATGTTTGAAGCTATTCTGACAGAAAAGACAAAGGTTACTAATTGTA
TGATAGTGTTTTTATATGGAAGATGTACAGCTTATGGACAAATGTACACCTTTTTGTTACTTTAATAAAAAATGT
AGTAGGATAAAAAAAA

589/6881
FIGURE 549

MSHGKGT DMLPEIAAAVGFLSSLLRTRGCVSEQRLKVFSGALQEALTEHYKHHWFPEKPSKSGYRCIRINH KMD
PIISRVASQIGLSQPQLHQLLPSELT LWVDPYEVSYRIGEDGSICVLYEEAPLAASCGLLTCKNQVLLGRSSPSK
NYVMAVSS

590/6881
FIGURE 550

ATGTCCATGAGGAGCCCCATCTCTGCCAGCTGGCCCTGGATGGCGTTGGCACCATGGTGAAGTGCACCATCAAG
TCAGAGGAGAAGAAAGAGCCTTGCCACGAGGCCCCCAGGGCTCAGCCACTGCCGCTGAACCTCAGCCTGGAGAC
CCAGCCCGGGCCTCCCAGGATAGTGCTGACCCCAAGCTCCAGCCCAGGGGAATTCAGGGGCTCCTGGGACTGT
AGCTCTCCAGAGGGTAATGGGTCCCCAGAACCAAGAGACCAGGAGCGTCGGAGGCTGCCTCTGGAAGCCAGGAG
AAGCTGGACTTCAACCGAAATTTGAAAGAAGTGGTGCCAGCCATAGAGAAGCTGTTGTCCAGTGAAGGAG
AGGTTTCTAGGAAGGAACCTATGGAAGCCAAAGATGTCAAAGGGACCCAAGAGAGCCTAGCAGAGAAGGAGCTC
CAGCTTCTGGTCATGATTACCCAGCTGTCCACCTGCGGGACAGCTCCTGACAGCCCACTCGGAGCAGAAGAAC
ATGGCTGCCATGCTGTTTGAGAAGCAGCAGCAGCAGATGGAGCTTGCCCGCAGCAGCAGGAGCAGATTGCAAAG
CAGCAGCAGCAGCTGATTACGACAGCAGCATAAGATCAACCTCCTTCAGCAGCAGATCCAGCAGTTAACATGCCCT
TATGTATGATCCAGCCTTCCCCCAAGCCACCAACCTCTGCCCTGTCACCCCTGACTCCAGCTGGCCTTACCC
ATTAGCCCATTCCTTGCAAACAGTGGAGTATCCGCTGCAGCTGCTGCACAGCCCCCTGCCCAAGTGGTGAAG
AGGCCTGGGGCCATGGCCACCCACACCCCTGCGAGGAGCCCTCCAGCCCTGAACCTCACAGCCAAGCCCAAG
GCCCCGAGCTGCCAACACCTCCAGCTCCCCAAGCCTGAAGATGAGCAGCTGTGTGCCCCGCCCCCCAGCCAT
GGAGGCCCCACGCGGGACCTGCAGTCCAGCCCCCGAGCCTGCCCTCTGGGCTTCTTGGTGAAGGGGACGCTGT
ACCAAAGCCATCCAGGATGCTCGGCAGCTGCTGCACAGCCACAGTGGGGCCTTGGATGGCTCCCCAACACCCCC
TTCCGTAAGGACCTCATCAGCCTGGACTCATCCCCAGCCAAGGAGCGGCTGGAGGACGGCTGTGTGCACCCACTG
GAGGAAGCCATGCTGAGCTGCGACATGGATGGCTCCCGCCACTTCCCGAGTCCCGAAACAGCAGCCACATCAAG
AGGCCCATGAACGCCCTTCATGGTGTGGGCCAAGGATGAGCGGAGGAAGATCCTGCAAGCCTTCCAGACATGCAC
AACTCCAGCATCAGCAAGATCCTTGATCTCGCTGGAAGTCCATGACCAACCAGGAGAAGCAGCCCTACTATGAG
GAACAGGCGCGGCTGAGCCGGCAGCACCTGGAGAAGTATCCTGACTACAAGTACAAGCCGCGGCCCAAGCGCACC
TGCATCGTGGAGGGCAAGCGGCTGCGCGTGGGAGAGTACAAGGCCCTGATGAGGACCCGCGCTCAGGATGCCCGC
CAGAGCTACGTGATCCCCCGCAGGCTGGCCAGGTGCAGATGAGCTCCTCAGATGTCCTGTACCCTCGGGCAGCA
GGCATGCCGCTGGCACAGCCACTGGTGGAGCACTATGTCCCTCGTAGCCTGGACCCCAACATGCCTGTGATCGTC
AACACCTGCAGCCTCAGAGAGGAGGGTGAGGGCAGAGATGACAGGCACCTCGGTGGCTGATGGCGAGATGTACCGG
TACAGCGAGGACGAGGACTCGGAGGGTGAAGAGAACAGCGATGGGGAGTTGGTGGTGTCTACAGACTGATCCCCG
CTGGGTGGGCCTGGCCCCCTTCTCCTCTGGGGAAAGACCTTGTCCCAACTCGATGGGCACAGCCAGCCAACCTAAGA
CTATGTTGGTACTTGGACTTGTTCGTGCCCCAGAGATGGGCAAAGCTGTGCACTTGAGATACATTATGAGGGG
AGAGGCGCCCTCCCTTCTGAGGAGCTGTTGGCCTGGGTGGGCAGGAAGTGCAGTATGGCCATGGGCTGAGCAGG
CTGAGCACCTCAGCCTTTAGGGCTTATGGCCAGGGGACACTGTATGACTCTCCTCTCCTGCAGGTGTCTATCCAC
CTGGGGTATGGCATCTACCGACCTGTCTCCCTGGGGTACATGCTTTGTTTCCATTCTGTCTGGCTGGACCAG
CCACTGTGGGACCAACACCCCTCCACACTCCCCAGACTGCTCGTCTATCACCAGGATCGCTTTGTACTTTGTG
CAAAAGGGTCTGGCTGTCCCTTGCTGTTTTCATCTCTGCCAAGCCTATTGTGCCCTCTGGCTGCTGTATGTGTGCG
CGTGACGCTGTGTGTTTTTCATCTGTTTCACTGCACAAGATATTTATTGAGTGGCCACTACGTGCCAGGCAC
TGTGTCTGAGTTCTGTGGGTGTGTCTCTCGATGCCACTCCTGCTTCTCTGGGGGCTCTTTCTGTGCTTCTCTT
TGTCCCCAAATTGCTACCTCTTTGTCTAGTCTGGGTGTCTCAGGTTCTGTGTGCTCTTGTGTGCATTTCTGTCTCT
CTCTGTCTCTGCTCTCTGCAAGGCCCTCTATTTCTCTCTTTCTTGGTGTCTGTCTTTGCCCCCTGTGCCCTCT
GGATTCTCTGGGTCTATGTAGGCCCTGGTCTGCCCTGGGCTCATCAGCCTTCTGACCTCCTCTGCCCTCCCC
TTCACTCCCTCCCTGGCTCTGCCAGTGGTTCCACGGAGCCATTTTTAGCTCTGATCAGCATGGGAATGTGCCT
CGGCCTCCAAGGGGCTTTGTCTGGTGGCCCCGCCCCCTGGTCCCAACCTGATCCACGAGGGAGTTGGGACAGGA
GGATTGATGGTGTCTCCCTTCTGCCAGCGTCAGAGGCCCTGGAGAGGGGCTGTCCATGGCAGCTGGTCTTTATT
CCTCCCTCATGAGCACAGGGTTCGGGGGGGTCCCCATTCTTGAAGAGGTTGAGAAGACTCCTGGGCTTCAGCCTC
TCCCACCCAGCCCTGCCCTCACCTGCCCTGCCCTCCCCCTCCCCACTCTATACTAGGGACTGGATCTCAGCCTCT
GATCAGTTTCAAAAGTTTGTTCCTAAGGAAATCAAATCCCATTGTACCTAACTCTGAAGATCTAAATAGCCC
TTGGATCAGTATGGGAACCCCAAATCCACAGGGCCAGATGTGGAGTCTGTGTCTGCCCCGCTCTCTCTCCATC
CTCAAAGCCCCCACTTCTCTCCAGGCTGTTTCTTTTTTATGACTGTAAACATAGATAGTCTTTATTTTGTAA
TAATAAGATAATGATGAGTAACCTAACCAGCACATTTCTCTGTTTACACTCGGGGGATTTTTTTGTTTTCTGAT
GACATAATAAAGACAGATCATTTCAGAAAAAAAAAAAAAAAAAAAA

591/6881
FIGURE 551

MSMRSPISAQLALDGVGTMVNCTIKSEEKKEPCHEAPQGSATAAEPQPGDPARASQDSADPQAPAQGNFRGSWDC
SSPEGNGSPEPKRPGASEAASGSQEKLDNFNRNLKEVVPKLLSSDWKERFLGRNSMEAKDVKGTOESLAEKEL
QLLVMIHQSTLRDQLLTAHSEQKNMAAMLFEKQQQOMELARQQEQIAKQQQQLIQQQHKINLLQQQIQQVNMP
YVMIPAFPPSHQPLPVTQDSQLALPIQPIPCPKVEYPLQLLHSPAPVVKRPGAMATHHPLQEPSQPLNLTAKPK
APELPNTSSSPSLKMSSCVPRPPSHGGPTRDLQSSPPLPLGFLGEGDAVTKAIQDARQLLHSHSGALDGSPTNP
FRKDLISLDSSPAKERLEDGCVHPLEEAMLSCDMDGSRHFPESRNSSHIKRPMAFMVWAKDERRKILQAFPMH
NSSISKILGSRWKSMTNQEKQPYEEQARLSRQHLEKYPDYKYKPRPKRTCIVEGKRLRVGEYKALMRTTRQDAR
QSYVIPQAGQVQMSSSDVLYPRAAGMPLAQPLVEHYVPRSLDPNMPVIVNTCSLREEGEGTDDRHSVADGEMYR
YSEDEDSEGEKSDGELVVLTD

592/6881

FIGURE 552

AACTGCCATCTTCCAGTAATTCGCCAAAATGACGAATACAAAGGGAAAGAGGAGAGGCACCCGATGTATGTTCTC
TAGGCCTTTTAGAAAACATGGAGTTGTTCCTTTGGCCACGTATATGCGAGTCTATAAGAAAGGTGATATTGTAGA
CATCAAGGGAATGGGTACTGTTCAAAAAGGAATGCCCCACACGCGTTACCACGGCAAACTGGAAGAGTCTACAA
TGTTACCCAGCATGCTGTTGGCATTGTTGTAAACAAACAAGTTAAAGGCAAGATTCTTGCCAAGAGAATTAATGT
GCGTATTGAGCACATTAAAGCACTCTAAGAGCCGAGATAGCTTCCTGAAACGTGTGAAGGAAAATGATCAGAAAAA
GAAAGAAGCCAAAGAGAAAAGTTACCTGGATTCAACTAAAGCGCCAGCCTGCTCCACCCAGAGAAGCACACTTTGT
GAGAACCAATGGGAAGGAGCCTGAGCTGCTGGAACCTATTCCCTATGAATTCATGGCATAATAGGTGTTAAAAA
AAAAA

593/6881
FIGURE 553A

GCTTTTCAGGCGATCTGGAGAAAGAACGGCAGAACACACAGCAAGGAAAGGTCCTTTCTGGGGATCACCCCATTGG
CTGAAGATGAGACCATTCTTCTCTTGTGTTTTGCCCTGCCCTGGCCTCCTGCATGCCCAACAAGCCTGCTCCCGT
GGGGCCTGCTATCCACCTGTTGGGGACCTGCTTGTGGGAGGACCCGGTTTCTCCGAGCTTCATCTACCTGTGGA
CTGACCAAGCCTGAGACCTACTGCACCCAGTATGGCGAGTGGCAGATGAAATGCTGCAAGTGTGACTCCAGGCAG
CCTCACAACTACTACAGTCACCGAGTAGAGAATGTGGCTTCATCCTCCGGCCCCATGCGCTGGTGGCAGTCCAG
AATGATGTGAACCCTGTCTCTCTGCAGCTGGACCTGGACAGGAGATTCCAGCTTCAAGAAGTCATGATGGAGTTC
CAGGGGGCCATGCCCGCCGGCATGCTGATTGAGCGCTCCTCAGACTTCGGTAAGACCTGGCGAGTGTACCACTAC
CTGGCTGCCGACTGCACCTCCACCTTCCCTCGGGTCCGCCAGGGTCCGGCTCAGAGCTGGCAGGATGTTCGGTGC
CAGTCCCTGCCCTCAGAGGCCTAATGCACGCCTAAATGGGGGGGAGGTCCAACCTTAACCTTATGGATTTAGTGTCT
GGGATTCCAGCAACTCAAAGTCAAAAATTCAAGAGGTGGGGGAGATCACAACTTGAGAGTCAATTTACCAGG
CTGGCCCCCTGTGCCCAAAGGGGCTACCACCCTCCAGCGCCTACTATGCTGTGTGCCAGCTCCGTCTGCAGGGG
AGCTGCTTCTGTACGGCCATGCTGATCGCTGCGCACCCAAGCCTGGGGCCTCTGCAGGCCCCCTCCACCGCTGTG
CAGGTCCACGATGTCTGTGTCTGCCAGCACAACTGCCGGCCCAAATTGTGAGCGCTGTGCACCCCTTCTACAAC
AACCGGCCCTGGAGACCGGGCGGAGGGCCAGGACGCCCATGAATGCCAAAGGTGCGACTGCAATGGGCACTCAGAG
ACATGTCACTTTGACCCCGCTGTGTTGCCGCCAGCCAGGGGGCATATGGAGGTGTGTGTGACAATTGCCGGGAC
CACACCGAAGGCAAGAACTGTGAGCGGTGTGAGTGCAGTACTATTTCCGGAACCGGCGCCCGGGAGCTTCCATTGAG
GAGACCTGCATCTCCTGCGAGTGTGATCCGGATGGGGCAGTGCAGGGGCTCCCTGTGACCCAGTGACCGGGCAG
TGTGTGTGCAAGGAGCATGTGCAGGGAGAGCGCTGTGACCTATGCAAGCCGGGCTTCACTGGACTCACCTACGCC
AACCCGCAGGGGTGCCACCGCTGTGACTGCAACATCCTGGGGTCCCGAGGGACATGCCGTGTGACGAGGAGAGT
GGGCGCTGCCTTTGTCTGCCAACGTGGTGGGTCCCAAATGTGACCACTGTGCTCCCTACCACTGGAAGCTGGCC
AGTGGCCAGGGCTGTGAACCGTGTGCTGCGACCCGACAACTCCCTCAGCCACAGTGCAACCAAGTTCACAGG
GCAGTGCCTGTGCGGAAGGCTTTGGTGGCCTGATGTGCAGCGCTGCAGCCATCCGCCAGTGTCCAGACCGGACC
TATGGAGACGTGGCCACAGGATGCCGAGCCTGTGACTGTGATTTCCGGGAACAGAGGGCCCGGGCTGCGACAAG
GCATCAGGCCGCTGCCTCTGCCGCCCTGGCTTGACCGGGCCCCGCTGTGACCACTGCCAGCGAGGCTACTGCAAT
CGCTACCCGGTGTGCGTGGCCTGCCACCCTTGCTTCCAGACCTATGATGCGGACCTCCGGGAGCAGGCCCTGCGC
TTTGGTAGACTCCGCAATGCCACCGCCAGCCTGTGGTCAGGGCCTGGGCTGGAGACCGTGGCCTGCCCTCCCGG
ATCCTAGATGCAAAGAGTAAGATTGAGCAGATCCGAGCAGTTCTCAGCAGCCCCGCAGTCACAGAGCAGGAGGTG
GCTCAGGTGGCCAGTGCCATCCTCTCCCTCAGGCGAACTCTCCAGGGCCTGCAGCTGGATCTGCCCTGGAGGAG
GAGACGTTGTCCCTTCCGAGAGACCTGGAGAGTCTTGACAGAAGCTTCAATGGTCTCCTTACTATGTATCAGAGG
AAGAGGGAGCAGTTTGA AAAAATAAGCAGTGTGATCCTCAGGAGCCTTCCGGATGCTGAGCACAGCCTACGAG
CAGTCAGCCCAGGCTGCTCAGCAGGTCTCCGACAGCTCGCGCCTTTTGGACCACTCAGGGACAGCCGGAGAGAG
GCAGAGAGGCTGGTGGCGCAGGCGGGAGGAGGAGGACCGGCAGCCCCAAGCTTGTGGCCCTGAGGCTGGAG
ATGCTCTCGTTGCTGACCTGACACCCACCTTCAACAAGCTCTGTGGCAACTCCAGGCAGATGGCTTGACCCCCA
ATATCATGCCCTGGTGAGCTATGTCCCAAGACAATGGCACAGCCTGTGGCTCCCGCTGCAGGGGTGTCTTCCC
AGGGCCGGTGGGGCCTTCTTGATGGCGGGCAGGTGGCTGAGCAGCTGCGGGGCTTCAATGCCAGCTCCAGCGG
ACCAGGCAGATGATTAGGGCAGCCGAGGAATCTGCCTCACAGATTCAATCCAGTGCCCAGCGCTTGGAGACCCAG
GTGAGCGCCAGCCGCTCCAGATGGAGGAAGATGTGAGACGCACACGGCTCCTAATCCAGCAGGTCCGGGACTTC
CTAACAGACCCCGACACTGATGCAGCCACTATCCAGGAGGTGAGCGAGGCGTGTGCTGGCCCTGTGGCTGCCACA
GACTCAGCTACTGTTCTGCAGAAGATGAATGAGATCCAGGCCATTGCAGCCAGGCTCCCCAACGTGGACTTGGTG
CTGTCCAGACCAAGCAGGACATTGCGCGTCCCCGCCGTTGACAGGCTGAGGCTGAGGAAGCCAGGAGCCGAGCC
CATGCACTGGAGGGCCAGGTGGAAGATGTGTTGGGAACCTGCGGCAGGGGACAGTGGCACTGCAGGAAGCTCAG
GACACCATGCAAGGCACCGCCTCCCTTCCGCTTATCCAGGACAGGGTTGCTGAGGTTTCCAGGTTACTGCGG
CCAGCAGAAAAGCTGGTGACAAGCATGACCAAGCAGCTGGGTGACTTCTGGACACGGATGGAGGAGCTCCGCCAC
CAAGCCCGGCAGCAGGGGGCAGAGGCAGTCCAGGCCAGCAGCTTGCAGGAAGGTGCCAGCGAGCAGGCATTGAGT
GCCCCAAGAGGGGATTTGAGAGAATAAAACAAAAGTATGCTGAGTTGAAGGACCGGTTGGGTGAGGTTCCATGCTG
GGTGAGCAGGGTGCCCGGATCCAGAGTGTGAAGACAGAGGCAGAGGAGCTGTTTGGGGAGACCATGGAGATGATG
GACAGGATGAAAGACATGGAGTTGGAGCTGTGCGGGGCAGCCAGGCCATCATGCTGCGCTCGGCGGACCTGACA
GGACTGGAGAAGCGTGTGGAGCAGATCCGTGACCACATCAATGGGCGCGTGTCTACTATGCCACCTGCAAGTGA

594/6881
FIGURE 553B

TGCTACAGCTTCCAGCCCGTTGCCCCACTCATCTGCCGCCTTTGCTTTTGGTTGGGGGCAGATTGGGTTGGAATG
CTTTCCATCTCCAGGAGACTTTCATGCAGCCTAAAGTACAGCCTGGACCACCCCTGGTGTGTAGCTAGTAAGATT
ACCCTGAGCTGCAGCTGAGCCTGAGCCAATGGGACAGTTACACTTGACAGACAAAGATGGTGGAGATTGGCATGC
CATTGAAACTAAGAGCTCTCAAGTCAAGGAAGCTGGGCTGGGCAGTATCCCCGCCTTTAGTTCTCCACTGGGGA
GGAATCCTGGACCAAGCACAAAACTTAACAAAAGTGATGTAAAAATGAAAAGCCAAATAAAAAATCTTTGG

595/6881
FIGURE 554

MRPFFLLCFALPGLLHAQQACSRGACYPPVGDLLVGRTRFLRASSTCGLTKPETYCTQYGEWQMKCKCDSRQPH
NYYSHRVENVASSSGPMRWWQSQNDVNPVSLQLDLDRRFQLQEVMMEFQGPMPAGMLIERSSDFGKTWRVYQYLA
ADCTSTFPVRVQRQRPQSWQDVRCQSLPQRPNARLNGGKVQLNMDLVSGIPATQSQKIQEVGEITNLRVNFTRLA
PVPQRGYHPPSAYYAVSQLRLQGSCFCHGHADRCAPKPGASAGPSTAVQVHDVCVCQHNTAGPNCERCAPFYNNR
PWRPAEGQDAHECQRCDNCNGHSETCHFDPAVFAASQGAYGGVCDNCRDHTGKNCERCQLHYFRNRRPGASIQET
CISCECDPDGAVPGAPCDPVTGQCVCKEHVQGERCDLCKPGFTGLTYANPQGCHRCDCNILGSRRDMPCDDEESGR
CLCLEPNVVGPKCDQCAPYHWKLASGQGCEPCACDPHNSPQPTVQPVHRAVPCREGFGGLMCSAAAIRQCPDRTYG
DVATGCRACDCDFRGTEGPGCDKASGRCLCRPGLTGPRCDQCQRGYCNRYPCVACHPCFQTYDADLREQALRFG
RLRNATASLWSGPGLEDRLASRILDAKSKIEQIRAVLSSPAVTEQEVAQVASAILSLRRTLQGLQLDLPLEEET
LSLPRDLESLSRFSNGLLTMYQRKREQFEKISSADPSGAFRMLSTAYEQSAQAAQVSDSSRLDQLRDSRREAE
RLVRQAGGGGGTGSPKLVALRLEMSSLPDLTPTFNKLCGNSRQMACTPISCPGELCPQDNGTACGSRCRGVLPRA
GGAFLMAGQVAEQLRGFNAQLQRTROMIRAAEESASQIQSSAQRLTQVSASRSQMEEDVRRTRLIIQQVRDFLT
DPDIDAATIQEVSEAVLALWLPTDSATVLQKMNEIQAIARLPNVDLVLSQTKQDIARARRLQAEAEARSRAHA
VEGQVEDVVGNLRQGTVALQEAQDTMQGTSRSLRLIQDRVAEVQVLRPAEKLVTSMKQLGDFWTRMEELRHQA
RQQGAEAVQAQQLAEGASEQALSAQEGFERIKQKYAELKDRLGQSSMLGEQGARIQSVKTEAEELFGETMEMMDR
MKDMELELLRGSQAIMLRSADLTGLEKRVEQIRDHINGRVLYATCK

596/6881
FIGURE 555

GCGGAGTCTCCAACTGGGAGAGCTGCAGCTGCCGAGAGGAGGAGAACGCTGAGGTCGGTCGGACCAACGGACGCG
CTGACCGCTGCCAACTGCAGCTCGCGCTGCCTCCTGCTCGCGCCGTGCCACTAAGGTCCTCCGCTCCGAGAG
CCCAGAGCCGAGATGGAAACGGTCCAGGAGCTGATCCCCCTGGCCAAGGAGATGATGGCCCAGAAGCGCAAGGGG
AAGATGGTGAAGCTGTACGTGCTGGGCAGCGTGCTGGCCCTCTTCGGCGTGGTGCTCGGCCTGATGGAGACTGTG
TGCAGCCCCCTTCACGGCCGCCAGACGTCTGCGGGACCAGGAGGCAGCCGTGGCGGAGCTGCAGGCCGCCCTGGAG
CGACAGGCTCTCCAGAAGCAAGCCCTGCAGGAGAAAGGCAAGCAGCAGGACACGGTCCTCGGCGGCCGGGCCCTG
TCCAACCGGCAGCACGCCTCCTAGGAACTGTGGGAGACCAGCGGAGTGGGAGGGAGACGCAGTAGACAGAGACAG
ACCGAGAAGGAAGGGAGAGACAGAGGGGGCGCGCACAGGAGCCTGACTCCGCTGGGAGAGTGCAGGAGCACGT
GCTGTTTTTTATTTGGACTTAACTTCAGAGAAACCGCTGACATCTAGAAGTACCTACCACAAGCATCCACCAA
GGAGTTTGGGATTGAGTTTGTGCTGTGCAGCACTGCATTGTGATGACATTTCCAACACTGTGTGAATTATCTA
AATGCGTCTACCATTTTGCAGTAGGGAGGAAGGATAAATGCTTTTTATGTTATTATTATTATTATTACATGAC
CACCATTTTGCATTTTGAATAAAAACTTTTATACCAAAAAAAAAAAAAAAAAA

597/6881
FIGURE 556

METVQELIPLAKEMMAQKRKGKMKLYVLGSLALFGVVLGLMETVCSPFTAARRLRDQEAAVAELQAALERQAL
QKQALQEKKGKQDDTVLGGRALSNRQHAS

598/6881
FIGURE 557

CCCGCCTCCGAGAGCCCAGAGCCGAGATGGAAACGGTCCAGGAGCTGATCCCCCTGGCCAAGGAGATGATGGCCC
AGAAGCGCAAGGGGAAGATGGTGAAGCTGTACGTGCTGGGCAGCGTGCTGGCCCTCTTCGGCGTGGTGCTCGGCC
TGATGGAGACTGTGTGCAGCGGAGTGGGAGGGAGACGCAGTAGACAGAGACAGACCGAGAGAGGAATGGAGAGAC
AGAGGGGGCGCGCGCACAGGAGCCTGACTCCGCTGGGAGAGTGCAGGAGCACGTGCTGTTTTTTATTGGACTTA
ACTTCAGAGA

599/6881
FIGURE 558

CCGCTGGGCGTAGCTGCGACTCGGCGGAGTCCCGGCGGCGCGTCCTTGTTCTAACCCGGCGCGCCATGACCGTCG
CGCGGCCGAGCGTGCCCGCGGCGCTGCCCCCTCCTCGGGGAGCTGCCCCGGCTGCTGCTGCTGGTGCTGTTGTGCC
TGCCGGCCGTGTGGGGTGAAGTGTGGCCTTCCCCCAGATGTACCTAATGCCAGCCAGCTTTGGAAGGCCGTACAA
GTTTTCCCGAGGATACTGTAATAACGTACAAATGTGAAGAAAGCTTTGTGAAAATTCCTGGCGAGAAGGACTCAG
TGATCTGCCTTAAGGGCAGTCAATGGTCAGATATTGAAGAGTTCTGCAATCGTAGCTGCGAGGTGCCAACAGGC
TAAATTCTGCATCCCTCAAACAGCCTTATATCACTCAGAATTATTTCCAGTCGGTACTGTTGTGGAATATGAGT
GCCGTCCAGGTTACAGAAGAGAACCTTCTCTATCACCAAAATACTTGCCTTCAGAATTTAAAATGGTCCACAG
CAGTCGAATTTTGTAAAAAGAAATCATGCCCTAATCCGGGAGAAATACGAAATGGTCAGATTGATGTACCAGGTG
GCATATTATTTGGTGCAACCATCTCCTTCTCATGTAACACAGGGTACAAATTATTTGGCTCGACTTCTAGTTTTT
GTCTTATTTTCAAGCAGCTCTGTCCAGTGGAGTGACCCGTTGCCAGAGTGACAGAGAAATTTATTGTCCAGCACCAC
CACAAATTGACAAATGAATAATCAAGGGGAACGTGACCATTATGGATATAGACAGTCTGTAACGTATGCATGTA
ATAAAGGATTACCATGATTGGAGAGCACTCTATTTATTGTACTGTGAATAATGATGAAGGAGAGTGGAGTGGCC
CACCACCTGAATGCAGAGGAAAATCTCTAAGTTCCAAAGTCCCACCAACAGTTCAGAAACCTACCACAGTAAATG
TTCCAACCTACAGAAGTCTCACCACCTTCTCAGAAAACCAACCAACCAACCAACCAACCAACCAACCAACCAAC
GGAGTACACCTGTTTCCAGGACAACCAAGCATTTCATGAAACAACCCCAAATAAAGGAAGTGGAACCACTTCAG
GTACTACCCGCTCTTCTATCTGGGCACACGTGTTTCACGTTGACAGGTTTGCTTGGGACGCTAGTAACCATGGGCT
TGCTGACTTAGCCAAAGAAGAGTTAAGAAGAAAATACACACAAGTATACAGACTGTTTCCTAGTTTCTTAGACTTA
TCTGCATATTGGATAAAAATAAATGCAATTGTGCTCTTCATTTAGGATGCTTTCATTGTCTTTAAGATGTGTTAGG
AATGTCAACAGAGCAAGGAGAAAAAGGCAGTCCTGGAATCACATTCTTAGCACACCTACACCTCTTGAAAATAG
AACAACCTTGCAAGATTGAGAGTGATTCTTTTCTTAAAGTGTAAGAAAGCATAGAGATTTGTTGCTATTTAGAA
GGGATCACGAGGAAAAGAGAAGGAAAGTGATTTTTTCCACAAGATCTGTAATGTTATTTCCACTTATAAAGGAA
ATAAAAAATGAAAAACATTATTTGGATATCAAAAGCAAATAAAAACCCAATTGAGTCTCTTCTAAGCAAAATTGC
TAAAGAGAGATGAACCACATTATAAAGTAATCTTTGGCTGTAAGGCATTTTCATCTTCTTCCGGGTGGCAAAA
TATTTTAAAGGTAAACATGCTGGTGAACCAAGGGGTGTTGATGGTGATAAGGGAGGAATATAGAATGAAAGACTG
AATCTTCTTTGTTGCACAAATAGAGTTTGAAAAAGCCTGTGAAAGGTGCTTCTTTGACTTAATGTCTTTAAA
AGTATCCAGAGATACTACAATATTAACATAAGAAAAGATTATATATTATTTCTGAATCGAGATGCCATAGTCAA
ATTTGTAAATCTTATCTTTTGTAAATATTATTTATATTTATGACAGTGAACATTCTGATTTTACATGTAA
AACAAGAAAAGTTGAAGAAGATATGTGAAGAAAATGTATTTTCTTAAATAGAAATAAATGATCCCATTTTTTG
GT

600/6881
FIGURE 559

MTVARPSVPAALPLLGE LPRLLLLVLLCLPAVWGDCGLPPDV PNAQPALEGRTSFPEDTVITYKCEESFVKIPGE
KDSVICLKGSQWSDIEEFCNRSCEVPTRLNSASLKQPYITQNYFPVGT VVEYECRPGYRREPSLSPKLTICLQNLK
WSTAVEFCKKKSCPNPGEIRNGQIDVPGGILFGATISFSCNTGYKLFGSTSSFCLISGSSVQWSDPLPECREIYC
PAPPQIDNGIIQGERDHYGYRQSVTYACNKGFTMIGEHSIYCTVNNDEGEWSGPPPECRGKSLTSKVPPTVQKPT
TVNVPTTEVSPTSQKTTTKTTTPNAQATRSTPVSRTTKHFHETTPNKGSGTTS GTTRLLSGHTCFTLTGLLGTLV
TMGLLT

601/6881
FIGURE 560A

GTGGATTGGTCGTCTCCCTGATTCCGAGCTGCGGGCAGGGAGAGGGGCCCTCGCGCCGCCCTCAGCAGCCGGCGG
CGGCCGAGGTAGACCGAGCGGGGACGGAAGGACAGACCGACGTCGCCGAGCTGGAATCATGTGAGGGCCAACCGG
GGAAGGTGGAGCAGATGAGCACACACAGGAGCCGTCTCCTCACCGCCGCCCTCTCAGCATGGAACAGAGGCGGC
CCTGGCCCCGGGCCCTGGAGGTGGACAGCCGCTCTGTGGTCTGCTCTCAGTGGTCTGGGTGCTGCTGGCCCCC
CAGCAGCCGGCATGCCTCAGTTCAGCACCTTCCACTCTGAGAATCGTGA CTGGACCTTCAACCACTTGACCGTCC
ACCAAGGGACGGGGGCCGTCTATGTGGGGGCCATCAACCGGGTCTATAAGCTGACAGGCAACCTGACCATCCAGG
TGGCTCATAAGACAGGGCCAGAAGAGGACAACAAGTCTTGTTACCCGCCCTCATCGTGACGCCCTGCAGCGAAG
TGCTCACCCTCACCACAATGTCAACAAGCTGCTCATCTGACTACTCTGAGAACC GCCTGCTGGCCTGTGGGA
GCCTCTACCAGGGGTCTGCAAGCTGCTGCGGCTGGATGACCTCTTCATCCTGGTGGAGCCATCCCAAGAAGG
AGCACTACCTGTCCAGTGTCAACAAGACGGGCACCATGTACGGGGTGATTGTGCGCTCTGAGGGTGAGGATGGCA
AGCTCTTCATCGGCACGGCTGTGGATGGGAAGCAGGATTACTTCCGACCTGTCCAGCCGGAAGCTGCCCCGAG
ACCTTGAGTCTCAGCCATGCTCGACTATGAGCTACACAGCGATTTTGTCTCCTCTCTCATCAAGATCCCTTCAG
ACACCCTGGCCCTGGTCTCCCACTTTGACATCTTCTACATCTACGGCTTTGCTAGTGGGGGCTTTGTCTACTTTC
TCACTGTCCAGCCCGAGACCCCTGAGGGTGTGGCCATCAACTCCGCTGGAGACCTCTTCTACACCTCAGCATCG
TGCGGCTCTGCAAGGATGACCCCAAGTTCCACTCATACGTGTCCCTGCCCTTCGGCTGCACCCGGGCGGGGTGG
AATACCGCCTCCTGCAGGCTGCTTACCTGGCCAAGCCTGGGGACTCACTGGCCAGGCCTTCAATATCACCAGCC
AGGACGATGTACTCTTTGCCATCTTCTCCAAAGGGCAGAAGCAGTATCACCACCCGCCGATGACTCTGCCCTGT
GTGCCTTCCCTATCCGGGCCATCAACTTGCAGATCAAGGAGCGCCTGCAGTCTTGCTACCAGGGCGAGGGCAACC
TGGAGCTCAACTGGCTGTGGGGAAGGACGTCCAGTGCACCAAGGCGCCTGTCCCATCGATGATAACTTCTGTG
GACTGGACATCAACCAGCCCTGGGAGGCTCAACTCCAGTGGAGGGCCTGACCCTGTACACCACCAGCAGGGACC
GCATGACCTCTGTGGCCTCCTACGTTTACAACGGCTACAGCGTGTTTTTGTGGGACTAAGAGTGGAAGCTGA
AAAAGATTCGGGCCGACGGTCCCCCCCATGGTGGGGTCCAGTACGAGATGGTCTCTGTGCTCAAGGACGGAAGCC
CCATCCTCCGGGACATGGCCTTCTCCATTGATCAGCGCTACCTGTACGTATGTCTGAGAGACAGGTACCAGGG
TCCCCGTGGAGTCATGTGAGCAGTATACGACTTGTGGGGAGTGCCTGAGCTCTGGGGACCCCTCACTGTGGCTGGT
GTGCCCTGCACAACATGTGCTCCCGCAGGGACAATGCCAACAGGCCTGGGAACCTAATCGATTTGTGTCAGCA
TCAGCCAGTGTGTGAGCCTTGCAGTGCATCCAGCAGCATCTCAGTATCTGAGCACAGCCGGTTGCTTAGCCTGG
TAGTGAGTGATGCTCCTGATCTATCTGCGGGTATCGCCTGTGCCTTTGGGAACCTGACAGAGGTGGAGGGGCAGG
TGTCCGGGAGCCAGGTATCTGCATCTCACCTGGGCCCCAAGGATGTCCCTGTCTATCCCGCTGGATCAAGACTGGT
TTGGGCTGGAGCTACAGCTGAGGTCCAAGGAGACAGGGAAGATATTGTGAGCACCGAGTTCAAGTTTTACAAC
GCAGTGGCCACCAACTGTGCCTGTCTGTGTCAACAGCGCCTTCCGCTGCCATTGGTGAAGTACCGCAACCTCT
GCACTCATGACCCACCACCTGCTCCTTCCAGGAGGCGGATCAATATTTAGAGGACTGTCCCCAGCTGGTG
CCACAGAGGAGATCTTGATTCCAGTCGGGGAGGTAAAGCCAAATCACCCTTAAGGCGCGAAATCTGCCCCAGCCGC
AGTCCGGCCAGCGAGGCTATGAGTGTGTCTCAACATACAAGGAGCCATCCACCGGGTCCCCGCTCTGCGCTTCA
ACAGCTCCAGCGTTCAAGTGTGAGAACAGCTCGTACCAGTATGATGGCATGGACATCAGCAATCTGGCCGTGGATT
TCGCTGTGGTGTGGAACGGCAATTTCATCATTGACAACCCTCAGGACCTGAAAGTCCATCTCTACAAGTGTGCAG
CCCAGCGGGAGAGCTGCGGCCCTCTGCCTCAAGGCCGACCGGAAGTTTGAGTGTGGCTGGTGCAGCGGCGAGCGCA
GGTGCACCCTCCACCAGCACTGTACCAGCCCTTCCAGCCCTGGCTCGACTGGTCCAGCCACAATGTCAAGTGTCT
CCAACCTCAAATCACCAGAGATTTGACGGTGTCTGGACCGCCGGAAGGAGGGACGCGAGTGACCATCCATGGCG
TGAACCTGGGTCTGGACTTCTCCGAGATCGCCACCATGTGCAGGTGGCTGGGGTGCCCTGCACGCCCCCTCCAG
GGGAATACATCATCGCTGAGCAGATTGTCTGTGAGATGGGCCATGCCCTCGTGGGAACCACCTCCGGGCCAGTAC
GCCGTGTGATTGGCGAGTGTAAGCCAGAGTTTATGACGAAGTCCCATCAGCAGTACACCTTCGTGAACCTTCTG
TGCTGTCACTCAACCCAATCCGAGTCCCGAGTCAGGAGGCACTATGGTGACCATTACCGGCCATTACCTTGGGG
CTGGGAGCAGCGTGGCAGTCTACCTGGGCAACCAGACCTGCGAGTTCTACGGGAGGTCAATGAGTGAGATCGTGT
GTGTCTACCCCCATCATCCAATGGCCTTGCCCCGTCCCTGTTTTCTGTGAGTGTGACCGAGCCCATGTGGATA
GCAACCTGCAGTTTGAGTACATAGATGACCCTCGGGTCCAGCGCATCGAGCCAGAGTGAGCATTGCCAGTGGCC
ACACACCCCTGACCATCACAGGCTTCAACCTGGATGTCAATCAGGAGCCAAGGATCCGAGTCAAATTCATGGCA
AAGAATCTGTCAATGTGTGTAAAGTTGTGAACACAACCACCTCACCTGCCTGGCACCCCTCTTGACCACGGACT
ACCGCCCTGGCCTGGACACTGTGGAACGCCCAGATGAGTTTGGATTTGTCTTTAACAATGTCCAATCCTTGCTAA

602/6881
FIGURE 560B

TTTACAACGACACCAAGTTTATCTACTACCCCAACCCGACCTTTGAACTGCTTAGCCCTACTGGAGTCTTGGATC
AAAAGCCAGGATCGCCCATCATTCTGAAGGGCAAAAACCTCTGCCCTCCTGCCTCTGGAGGGGCCAAACTCAACT
ACACTGTGCTCATCGGAGAGACCCCTTGTGCTGTACCGTATCTGAGACCCAGCTTCTCTGCGAGCCTCCCAACC
TCACCGGGCAGCACAAGGTCATGGTTACGTGGGCGGGATGGTGTCTCGCCTGGCTCGGTGAGTGTATCTCAG
ACAGCTTGTGACCCTGCCAGCCATCGTTCAGCATCGCGGGCCGGCGGCAGCCTCCTCCTCATCATCGTCATCATCG
TCCTCATTCCTTACAAGCGCAAGTCTCGAGAAAATGACCTCACTCTCAAGCGGCTGCAAATGCAGATGGACAATC
TGGAGTCCCGTGTGGCCTTGGAGTGCAAGGAAGCTTTTGTGAGCTCCAGACGGATATCAATGAGTTGACCAGTG
ACCTGGACCGCTCAGGAATCCCTTACCTGGACTATCGTACCTACGCTATGCGAGTCTGTTCCTGGGCATCGAGG
ACCACCCCGTCTGCGGGAGCTGGAGGTACAAGGAAACGGGCAGCAGCAGTGGAGAAGGCCCTGAAGCTCTTTG
CCCAGCTCATCAACAACAAGGTGTTCTGTGACCTTCATCCGCACCCTGGAGCTGCAGCGCAGTTTCTCCATGC
GCGACCGGGGCAACGTGGCTTCGCTCATCATGACCGGCTGCAGGGCCGCTGGAATATGCCACTGATGTCTCA
AGCAGCTGCTCTCTGACCTCATCGATAAGAACCTGGAGAACAAGAACCCCAAGCTGCTACTCCGGAGGACAG
AGTCTGTGGCTGAAAAGATGCTGACCAATTGGTTTCGCTTCTCTGACACAAGTTCCTAAAGGAGTGCGCAGGGG
AGCCACTCTTCATGCTATACTGTGCCATCAAGCAGCAGATGGAGAAGGGCCCCATTGATGCCATCACGGGCGAGG
CCCGCTACTCCCTGAGCGAGGACAAGCTCATCCGGCAGCAGATCGAGTACAAGACCCTGATCCTGAACTGCGTCA
ACCCTGACAACGAGAACAGTCCAGAGATCCAGTGAAGGTGTTAACTGTGACACCATCACACAGGTCAAGGAGA
AGATTCTTGATGCCGTGTATAAGAAATGTGCCCTATTCCAGCGGCCGAGGGCAGTGGACATGGACTTGGAGTGGC
GCCAAGGCCGGATCGCCCGGGTCGTGCTGCAAGATGAGGACATCACCACCAAGATTGAGGGTGACTGGAAGCGGC
TCAACACACTGATGCATTATCAGGTGTCAGACAGGTGCTGGTGGTGGCTCTGGTCCCCAAACAGACCTCCTCTACA
ACATCCCTGCCTCTGCCAGCATCTCCCGGACGTCCATCAGCAGATACGGTGACTCCTCCTTCAGGTATACGGGCA
GCCCCGACAGCCTGCGGTCCCGGGCCCCGATGATACCCAGACCTGGAAAGTGGGGTCAAGGTGTGGCATCTGG
TGAAGAACCATGACCACGGTGACCAGAAGGAGGGTGACCGGGGCAGCAAGATGGTGTCCGAGATCTACCTGACCC
GGCTACTGGCCACCAAGGGCACCTGCAGAAGTTTGTGGACGACTTGTGAGACCTTGTTCAGCACTGTGCACC
GGGGCAGCGCTCTCCCCCTGGCCATCAAGTACATGTTTGATTTCCTAGATGAGCAGGCAGACAGGCACAGCATCC
ATGACACAGATGTGCGGCACACCTGGAAAAGCAACTGCCCTCCTGCGCTTCTGGGTGAACGTGATTAAGAACC
CCCAGTTCTGTGTTTGACATCCACAAGGGCAGCATCAGGACGCCTGCCTCTCTGTGGTGGCCAGACCTTCATGG
ACTCTTGTTCACGTGAGACACCGGTGGGCAAGGACTCCCCCTCCAACAAGCTGCTCTATGCCAAGGACATCC
CCAGCTACAAGAGCTGGGTGGAGAGATACTACGCAGACATCGCCAAGCTCCCAGCCATCAGTGACCAGGACATGA
ATGCCCTACCTCGCCGAGCAGTCCCGCTGCACGCCGTGGAGTTCAACATGCTGAGTGCCCTCAATGAGATCTACT
CCTATGTGCAAGTATAGTGAGGAGCTCATCGGGGCCCTAGAGCAGGATGAGCAGGCACGGCGGCAGCGGCTGG
CTTATAAGGTGGAGCAGCTCATTAAATGCCATGTCCATTGAGAGCTGAGAGGAGGAGCCTCGCATTCTGGGAAGA
GGGACCTGTCCAAGCTGTCACTGGGAGTCTCAGATGGAAGGACAAGTGATGGGGATCAGGCCCCAGAGCTTGC
TGTCCCTGAGACCCCATCCTGGGGAGAGGGGAGGACTCCTCTCCCTACGCCAGCCAAGTTTCGTATAGCCAGT
TCCAGCTGGGAGAGACAGTGGGCGTCTGTCATCTCAGTGAGAACACCAGAGAACCCGGGGCCGGGAGAAGGTGG
TTCTTCAAGCCGAGAGGCACGAGCTGGGGACAGTCTGCCTCTGTGACTGCTGCTTTGCATGAAACTCATTGTA
TGTATATTGGGGAAATAATGAGAACTTTATTTAATTTTTTAAGAAAAAGGGAAAAAACAGAAATAAAACAAAA
AGCCGCCCTGTTAATCCAAATTAAAAA

603/6881

FIGURE 561

ACTGAATTCAACAATTATTTATTGAGTGTCTATTATGTGCTAGATACTGAGACACATCAGAGAACAAAACAAAA
GCCCTGCCCTCGTCGGGCTTACAGTCTAGCACTTACTGCCAGTTAACCTGGAGGCTACCTGGAACCCCGGGCAAG
TCACCGCACCTCTGTGCCTCGGTCTCAGCTGCCTAATGGGAGAATAAGCAGACCTGGCTCAGACATGAATCATG
TGCTTGGTGTACTGCAGATGCCAACTGCATCCCCATAACCCACCCAGACAGTAGATAGGGCTGGAAGTTGA
TTTTTAATGATCAAGTACAATGGAGGGAGGGCAGAGGGGCTAAGCCTAGCTGTCTGGGGTGCTGTGGTGGTGGTG
GGCTGGCTACACAACTGTTGCTGCTGCTGCTGCTTCTTGGTGGCTGCCTTGCTGGCGAGGTCCTTGGCCTTCTC
TGTAAGCTGCCAGTGTGTCTCCTTTGCCTTCTCCTTGGCTTCCTTGGCTGTCTCAACAAGTGTTTTGAAGGGGC
CTCGCCTTGCAGCTTTGCCAAGATATATTCAAACCCCTTCATAGTCTTGGTCACATTGCTTTTGAACCAGGCAAG
ACCAAATTCCTGGACAGCTCTGGAGACACCAATAAGCTAGAGGAGACCCAGGCTTCCCGGCGGATTTCGGTCCG
GCCACTGTTGTCAGAGTTCACACAGTAAACACATCGTTTCTCCACCACC

604/6881
FIGURE 562

TTTCGGCCGGAACCGCCATCTTCCAGTAATTCGCCAAAATGACAAACACAAAGGGAAAGAGGAGAGGCACCCGAT
ATATGTTCTCTAGGCCTTTTAGAAAACATGGAGTTGTTTCCTTTGGCCACATATATGCGAATCTATAAGAAAGGTG
ATATTGTAGACATCAAGGGAACGGGTACTGTTCAAAAAGGAATGCCCCACAAGTGTACCATGGCAAAACTGGAA
GAGTCTACAATGTTACCCAGCATGCTGTTGGCATTGTTGTAAACAAACAAGTTAAGGGCAAGATTCTTGCCAAGA
GAATTAATGTGCGTATTGAGCACATTAAGCATTCTAAGAGCCGAGATAGCTTCCTGAAACGTGTGAAGGAAAATG
ATCCTCCACCCAGAGAAGCACACTTTGTGAGAACCAATGGGAAGGAGCCTGAGCTGCTGGAACCTATTCCCTATG
AATTCATGGCATAATAGGTGTT

605/6881

FIGURE 563

MTNTKGKRRGTQYMF SRPFRKHGVVALATYMQIYKKGDIVDIKGMGTVQKGMPHKCYHGKTGRVYNVPQHAVGIV
VNTQVKGKILAKRINVRIEHIKHSKS

606/6881
FIGURE 564

ATGAAGCTCTTATCTTTGGTGGCTGTGGTCGGGTGTTTGCTGGTGCCCCAGCTGAAGCCAACAAGAGTTCTGAA
GATATCCGGTGC AAATGCATCTGTCCACCTTATAGAAACATCAGTGGGCACATTTACAACCAGAATGTATCCCAG
AAGGACTGCAACTGCCTGCACGTGGTGGAGCCCATGCCAGTGCCCTGGCCATGACGTGGAGGCCTACTGCCTGCTG
TGGGAGTGCAGGTACGAGGAGCGCAGCACCAACCACCATCAAGGTCATCATTGTCATCTACCTGTCCGTGGTGGGT
GCCCTGTTGCTCTACATGGCCTTCCTGATGCTGGTGGACCCTCTGATCCGAAAGCCGGATGCATATACTGAGCAA
CTGCACAATGAGGAGGAGAATGAGGATGCTCGCTCTATGGCAGCAGCTGCTGCATCCCTCGGGGGACCCGAGCA
AACACAGTCCTGGAGCGTGTGGAAGGTGCCAGCAGCGGTGGAAGCTGCAGGTGCAGGAGCAGCGGAAGACAGTC
TTCGATCGGCACAAGATGCTCAGCTTAG

607/6881
FIGURE 565

MKLLSLVAVVGCLLVPPAEANKSSDIRCKCICPPYRNISGHIYNQNVSQKDCNCLHVVEPMFVPGHDVEAYCLL
CECRYEERSTTTIKVIVIVYLSVVGALLLYMAFLMLVDPLIRKPDAYTEQLHNEEENEDARSMAAAAASLGGPRA
NTVLERVEGAQQRWKLQVQEQRKTVFD RHKMLS

608/6881
FIGURE 566

ACCTGCACTGACTTTTTTCTCCTTTTGGAGGGAGAGCAGAGACCATGTCTGACATAGAAGAGGTGGTGAAGAGT
ACGAGGAGGAGGAGCAGGAAGAAGCAGCTGTTGAAGAGCAGGAGGAGGCAGCGGAAGAGGATGCTGAAGCAGAGG
CTGAGACCGAGGAGACCAGGGCAGAAGAAGATGAAGAAGAAGAGGAAGCAAAGGAGGCTGAAGATGGCCCAATGG
AGGAGTCCAAACCAAAGCCCAGGTCGTTTCATGCCCAACTTGGTGCTCCCAAGATCCCCGATGGAGAGAGAGTGG
ACTTTGATGACATCCACCGGAAGCGCATGGAGAAGGACCTGAATGAGTTGCAGGCGCTGATCGAGGCTCACTTTG
AGAACAGGAAGAAAGAGGAGGAGGAGCTCGTTTCTCTCAAAGACAGGATCGAGAGACGTCGGGCAGAGCGGGCCG
AGCAGCAGCGCATCCGGAATGAGCGGGAGAAGGAGCGGCAGAACCGCCTGGCTGAAGAGAGGGGCTCGACGAGAGG
AGGAGGAGAACAGGAGGAAGGCTGAGGATGAGGCCCGGAAGAAGAAGGCTTTGTCCAACATGATGCATTTTGGGG
GTTACATCCAGAAGXXXXXXXXXACAGAGCGGAAAAGTGGGAAGAGGCAGACTGAGCGGGAAAAGAAGAAGA
TTCTGGCTGAGAGGAGGAAGGTGCTGGCCATTGACCACCTGAATGAAGATCAGCTGAGGGAGAAGGCCAAGGAGC
TGTGGCAGAGCATCTATAACTTGGAGGCAGAGAAGTTCGACCTGCAGGAGAAGTTCAAGCAGCAGAAATATGAGA
TCAATGTTCTCCGAAACAGGATCAACGATAACCAGAAAGTCTCCAAGACCCGCGGGAAGGCTAAAGTCACCGGGC
GCTGGAAATAGAGCCTGGCCTCCTTCACCAAAGATCTGCTCCTCGCTCGCACCTGCCTCCGGCCTGCACTCCCC
AGTTCCTGGGCCCTCCTGGGCACCCAGGCAGCTCCTGTTTGGAAATGGGGAGCTGGCCTAGGTGGGAGCCACCA
CTCCTGCCTGCCCCACACCCACTCCACACCAGTAATAAAAAGCCACCACACACTG

609/6881
FIGURE 567

CCAGCCTTGCCCGGCAGAGGACTCTGGAGGATGAGGAGGAACAGGAGCGCGGAXGGCGGCGGCACCGCAACCTG
AGCTCCACCACGGACGATGAGGCTCCCAGGCTCAGCAGAATGGAGACCGGCAGCCTCTGCTTCTGAGAGACTACC
GAGCGTGGAAGAAGCAGAGGTGCCCAAGCCACTGCCCCAGCCTCCAAAGATGAGGACGAGGACATCCAGAGCAT
CCTCAGAACACGGCAGGAGCGGAGGCAGAGGCGGCAGGTGGTGGAGGCTGCACAGGCCCCATCCAGGAGAGGCT
GGAGGCAGAGGAGGGAGGAACAGCTTGAGCCCTGTGCAGGCCACACAGAAACCCCTAGTCTCCAAGAAGGAAC
GGAAATCCCACCTCGCCGAGACTGAGTCGGGAACAGCGGGCCCCCTGCGCTGGAGGAGGAGAGCTTGGTGGGCA
GGGAGCCAGAAGAGAGGAAGAAAGGGGTTCCAGAAAAGTCCCCAGTCTTGAGAGAAATCCTCCATGCCAAAGAAGA
CGGCACCTGAAAAGAGCCTGGTCTCCGATAAAACCTCCATCTCTGAGAAGGTGCTGGCCTCAGAGAAGACATCTC
TATCAGAGAAGATAGCAGTGTGAGAGAAAAGAAACAGCTCAGAGAAGAAGTCTGTTCTAGAAAAAACAGTGTCT
CTGAGAAGTCGCTGGCCCCAGGGATGGCACTGGGCTCAGGAAGGAGGCTGGTGTCTGAGAAAGCTTCCATCTTTG
AGAAGGCACTGGCCTCAGAGAAGAGCCCAACTGCAGATGCTAAGCCGGCCCCAAGAGGCCACAGCCTCAGAGCA
GCCCCTGCGCAGGAGCCGCCAGCCTCTGGGGGAAGCCAGCCACCACCAAGGAGCAGAGAGGAAGGGCCCTCCC
TGGGAAGAACCTGCCCTCTTTGGCAGAGCAGGGGGCTTCAGACCCTCCGACTGTGGCCTCCCGCCTCCCACCCGT
CACACTCCAGGTGAAAATCCCCAGCAAGGAGGAAGAGGCAGATATGTCTCACCCACACAGCGAACCTACAGCAG
CTCCCTCAAACGCTCCAGCCCCAGGACCATCTCCTTTTCGGATGAAACCCAAGAAAGAAAACCTCGGAAACAACCT
AACTCGCAGTGCCAGCATGAAGCTCCCAGACAACACAGTGAAGTTGGGAGAGAAGCTGGAGAGATACCACACGGC
CATACGAGATCAGAACTGTCAAGTCTCGGGGTCTGCCTTGCACTGAGTTATTCGTGGCTCCTGTGGGTGTAGC
CAGCAAGCGCCACCTCTTTGAGAAGGAACCTGGCGGGCCAGAGCCGAGCAGAACCAGCCTCCAGCCGGAAGGAGAA
CTTGAGGCTCTCAGGGGTGTGACATCAAGGCTCAACCTGTGGATCAGCAGGACCCAGGAATCTGGAGATCAGGA
CCCCCAGGAGGCACAGAAAGCATCATCTGCAACCGAGAGGACTCAGTGGGGACAGAAATCTGACTCCTCGCTGGA
CGCTGAGGTGTGACAAGCCCCGCCAAGACAGACCTGCAAGTCTTCGTCTCAAGGGACCTCCCTCATGCCAGGCCC
CTGCCCTTCACAGCAGCACCCCTTTCTCTCATTTGTCCCTGTTCCCTTTTTGCCTGTGGATCTGTTTGGCCAGGGT
CCCTGGGGTCAGGAATATTTGCAAGACTCAGCCAGCTCCTTCCCAGCCAGCCTCTTGGGGCTGGGACTTTCTCA
CCCTGCGGCAGGCACAACAGATGCTGGGACCCAGTCTCTGCCAGGTACAGCACAAGTGCACATCAGCACTATG
GGGCTATGTCTGCCCAGAGACCTCTGCTCCTTCTGCTCACATCCACAGTCAGGGCACGGCGCCCTCAAGAA
CTCCAGAGTCACCTGTCTCATCGGCTCCAGCAAGTGCCCTTTGTCTATGATGTCCCCCTTCTCTGAGGCCTGG
ACCCACCCATCTTTGTCCCTGGGGCTGCTCCAGCCACTGAGGCCCGCTCTGGCCAGGGGAGAAGGAGCTGCCG
TGCGTCTTCCCTGTGCCCCGTCTCCCTGCTTGGTTCTCCCTCCCTTCCCTGGCCGGCTGCCATGGCCAGGAGCT
AAGTGCCTTTTTGTGTGCAACCACTTACCCCTTCTCTGAAAACCTGTTCTCAGGAAGGATCTGATAAACTCATT
TACTCTCAGGTGTAAGAGACTGATGAGACCTTAGAAGCGAATTCCTCTCTGGAGGCCCTTGCTTTCTAGCAGAGTC
ACCTGAAGTGTGTGAGGAGGATCATATTTTCTCATCCCCCTCTTCTCACATTAAGGTGGTGGCTTGCCACTC
AGCAGTCTAGCTTGGTGACTGGGAAGTCCACATACAGGGCCAGGCCTACCCCTCTTCCCCACAAGCCCCCTCC
AACCCCCACCCCATGCTCTGGACCTCATGGCTCCTATGAGCTTGGAGCATGGTGAACCATCAGAGAATCTAGAA
CCAACCAAGCTAGGAACATCAGCCTGGTGCCTGTAAACCCCTTAAAGCTGTGGTTTACAACCTTTTCAAAAATTTA
AATCATTAG

611/6881
FIGURE 569

ATGCGCGGCGGCGGCGGCGAGCTGGTGCGGCCTCCGGGCTGCCGGGTCCAGTGGCACAAGGATTAAAGGAAGCGTTA
GTGGATACGCTCACCGGGATCCTATCCCCAGTACAGGAGGTGCGGGCGGGCTGCTGAAGAACAGATTAAGGTGCTG
GAGGTGACGGAGGAATTTGGTGTTCACTTGGCAGAACTGACTGTAGATCCCCAGGGGGCACTGGCAATCCGTACG
CTGGCATCAGTCATCTTGAAACAATATGTGGAGACTCACTGGTGTGCCCAATCAGAGAAATTTAGGCCTCCTGAA
ACTACAGAAAGGGCAAAAATTGTTATCCGGGAGCTATTGCCAATGGGTTGAGAGAATCGATAAGCAAAGTGCGC
TCCAGTGTGGCCTATGCAGTGTGAGCCATTGCCCACTGGGACTGGCCTGAAGCTTGGCCCCAACTCTTCAACCTG
CTCATGGAGATGTTGGTGAGCGGAGACTTAAATGCCGTCCATGGAGCCATGCGTGTGCTGACAGAATTCACCTCGT
GAAGTTACAGACACACAGATGCCACTTGTGTCTCTGTCATTCTCCCAGAGAIGTATAAGATCTTCAACCATGGCT
GAGGTGTATGGTATTCGAACCCGTTCCCGAGCCGTGGAGATTTTACCACCTGTGCCCATATGATCTGTAACATG
GAGGAGCTGGAAGGGTGACGCCAAAGTCCTGATCTTTCCCGTGGTACAGCAGTTCACAGAGGCCTTTGTTTACG
GCCCCCAGATACCAGATGGCCCCACATCTGACAGTGGGTTAAGATGGAGGTCTAAAGGCAGTGACAGCCCTA
GTGAAAACTTCCCAAAGCACATGGTGTCTCCATGCAGCAGATTCTGCCTATTGTTTGAACACCCTAACCAG
AGTGCAGCTTTTTATGTGAGGACAGAAGTAAATTACACAGAAGAAGTAGAAGATCCTGTGGATTCTGATGGTGAA
GTCTGGGCTTTGAAAATCTCGTCTTTAGCATTTTTGAATTTGTCCATGCTCTACTAGAAAATAGCAAATTCAAA
AGCACTGTTAAGAAAGCCTTGCCCTGAATTGATTTATTATATTATCCTGTACATGCAAATCACTGAGGAGCAGATT
AAAGTATGGACAGCCAAACCCCAACAATTTGTAGAAGATGAAGATGATGATACATTCTCCTATACTGTTAGAATA
GCAGCTCAAGACTTGTGTCTGGCTGTGGCCACAGATTCCAGAATGAAAGTGCAGCAGCCCTGGCTGCTGCAGCC
ACTCGACATTTACAAGAAGCTGAGCAAACCAAAAACAGTGGCACTGAGCACTGGTGGAAAGATCCATGAGGCATGC
ATGCTTGCCCTAGGCTCAGTGAAGGCCATCATCACTGACAGTGTGAAAAATGGCAGGATTCAATTTGACATGCAT
GGGTTCTTGACCAATGTCATCCTTGACAGACCTCAACCTCTCAGTGTCTCCTTTCTCTTGGGCCGGGCACTTTGG
GCTGCCAGTGGTTCACTGTGTCTATGTCCCTGAACTGATCCAGCAGTTCCTACAGGCAACAGTTAGTGGTCTT
CAGGAGACACAGCCCCATCAGTTCGAATTTCTGCAGTGAGAGCCATCTGGGGTTATTGTGACCAACTGAAAGTC
TCAGAGAGTACCCACGTGCTCCAGCCCTTCTCCAGCATCCTTGATGGCTTAATTCACCTAGCAGCCAGTTC
AGCTCAGAGGTCTCAACCTGGTGATGGAGACCCTGTGCATCGTTTGTACAGTAGACCCCGAATTCACAGCAAGC
ATGGAAGCAAATCTGCCCTTCACCATCGCCATTTTCTAAAGTACAGTAATGATCCCGTCGTCGCCTCACTG
GCTCAGGACATCTTCAAGGAGCTGTCCAGATTGAAGCCTGTCAGGGCCCAATGCAAATGAGGCTGATTCCCACT
CTGGTCAGCATAATGCAGGCCCCAGCAGACAAGATTCTGCAGGGCTTTGTGCGACAGCCATTGATATCCTGACA
ACAGTAGTACGAAATACAAAGCCTCCCTTTCCAGCTTCTCATCTGCCAAGCTTTCCCTGCTGTGGCACAGTGT
ACCCTTCACACAGATGACAATGCCACCATGCAGAATGGCGGAGAGTGCTTGCGGGCCTATGTGTCACTGACCCTG
GAACAAGTAGCCAGTGGCATGATGAGCAGGGCCACAATGGACTGTGGTATGTGATGCAAGTGGTGAGCCAGCTC
CTGGACCCCGCACCTCAGAGTTCACTGCGGCCTTTGTGGGCCGCTTGTTCACCCCTCATCTCCAAGGCAGGG
CGGGAACCTCGGGGAGAATCTAGACCAGATTCTTCGTGCCATCCTCAGTAAGATGCAGCAGGCAGAGACGCTCAGT
GTCATGCAGTCCCTGATCATGGTGTTTCGCTCATCTGGTGACACTCAGCTAGAACCTCTCTTGGAGTTCTCTGTGT
AGCCTCCCAGGACCTACTGGCAAACCTGCTCTAGAGTTTGTGATGGCTGAGTGGACAAGCCGACAGCACCTGTTT
TATGGACAGTATGAAGGCAAAGTCAGCTCTGTGGCACTCTGTAAGCTGCTCCAGCATGGCATCAATGCAGATGAC
AAACGGCTACAGGATATCCGTGTGAAGGGAGAGGAGATCTACAGCATGGATGAGGGCATCCGCACCCGCTCTAAG
TCAGCCAAAACCCAGAACGCTGGACAAACATTCTTTGCTGGTCAAGATCCTAAAGCTGATCATCAACGAGCTC
TCCAACGTCTAGGAGCTAATGCCGCTCGCCAGGCCACTCCTGCAGAGTGGAGTCAAGATGACTCCAATGATATG
TGGGAGGACCAGGAGGAGGAAGAGGAGGAGGAGGATGGTTTAGCTGGCCAACCTTTATCTGACATTCTTGCT
ACAAGTAAATATGAGGAGGATTACTACGAGGATGATGAGGAAGATGACCCTGATGCCCTGAAGGATCCTCTCTAT
CAGATTGATCTGCAGGCATATCTCACAGATTTCCTCTGCCAGTTTGTCTCAGCAGCCCTGCTACATAATGTTTTCA
GGCCACCTTAATGACAATGAGAGGCGAGTTCTACAGACCATCGGCATCTAA

612/6881
FIGURE 570

MAAAAAAGAASGLPGPVAQGLKEALVDILTGLSPVQEVRAAAEEQIKVLEVTEEFVHLELTVDPPQALAIRQ
LASVILKQYVETHWCAQSEKFRPPETTERAKIVIRELLPGLRESISKVRSSVAYAVSAIAHWDWPEAWPQLFNL
LMEMLVSGDLNAVHGAMRVLTEFTREVTDITQMLVAPVILPEMYKIFTMAEVYGIRTRSRAVEIFTTCAHMICNM
EELEKGAAKVLIFPVVQQFTEAFVQALQIPDGPTSDSGFKMEVLKAVTALVKNFPHMVSSMQQILPIVWNTLTE
SAAFYVRTEVNYTEEVDPVSDGEVLGFENLVFSIFEVHALLSKFKSTVKKALPELIYYIILYMQITEEQI
KVWTANPQQFVEDEDDDTFSYTVRIAAQDLLLLAVATDFQNESAAALAAAATRLQEAQTKNSGTEHWWKIHEAC
MLALGSVKAIITDSVKNGRIHFDMHGFLTINVILADNLVSPFLLGRALWAASRFTVAMSPELIQQFLQATVSGL
HETQPPSVRISAVRAIWGYCDQLKVSESTHVLQPFPLSILDGLIHLAAQFSSEVLNLVMTLCIVCTVDPEFTAS
MESKICPFTTIAIFLKYSNDPVVASLAQDIFKELSQIEACQGPQMRLIPTLVSIMQAPADKIPAGLCATAIDILT
TVVRNTKPPLSQLLICQAFPAVAQCTLHTDDNATMQNGGECLRAYVSVTLEQVAQWHDEQGHNLWYVMQVVSQ
LDPRTEFTAAAFVGRVSTLISKAGRELGENLDQILRAILSKMQQAETLSVMQSLIMVFAHLVHTQLEPLLEFLC
SLPGPTGKPALEFVMAEWTSRQHLYGQYEGKVSSVALCKLLQHGINADDKRLQDIRVKGEEIYSMDGIRTRSK
SAKNPERWTNIPLLVKILKLIINELSNVMEANAARQATPAEWSQDDSDNMWEDQEEEEEEEDGLAGQLLSDILA
TSKYEEDYEDDEEDDPDALKDPLYQIDLQAYLTDFLCQFAQQPCYIMFSGHLNDNERRVLQTIGI

613/6881
FIGURE 571

AGTCAAGATGGAGGAGTACGCGGAGAGCCTTGCCCATGGCGAATTGTGGATGACTGTGGTGGGGCCTTTACGAT
GGGTACCATTGGTGGTGGTATCTTTCAAGCAATCAAAGGTTTTTCGCAATTCTCCAGTGGGAGTAAACCACAGACT
ACGAGGGAGTTTGACAGCTATTAAAACCAGGGCTCCACAGTTAGGAGGTAGCTTTGCAGTTTGGGGAGGGCTGTT
TTCCATGATTGACTGTAGTATGGTTCAAGTCAGAGGAAAGGAAGATCCCTGGAACTCCATCACAAGTGGTGCCTT
AACGGGAGCCATACTGGCAGCAAGAAATGGACCAGTGGCCATGGTTGGGTGAGCCGCAATGGGTGGCATTCTCCT
AGCTTTAATTGAAGGAGCTGGTATCTTGTGACAAGATTTGCCTCTGCACAGTTTCCCAATGGTCCCTCAGTTTGC
AGAAGACCCCTCCAGTTGCCTTCAACTCAGTTACCTTCCTCACCTTTTGGAGACTATCGACAATATCAGTAGGA
CTTCTTTCCTAGGATTTCTTTAACAGAACGAGTTGTGGTTCGAGAAGGATTTCAGAAGATCAAGTTACAGTCTGT
TTTTAAAACCATAGGTGGGACAGCTATGGCCAATAGGCTATAAAGAGACATTTAGCACTTTTTTCTATTTAAAGG
AACAAGCGGGGAAGGGTGCTAAAAGATAATACGTTTATTTATTCACACTTGAATTGCATTTGTGATCAAAATAAA
TGTTTAAATCGCTAAAGGAAAATACAGTAAGTGCTTGAAAGATGAAGGACCAAAAGGCCAAAAACAGTGAAATA
TGATCATCATCTCTTGCGGACTTCTCTGCCTGGTTTTGTGTGTTCTGTTATTCAAACAATAAAAGCTGGTGGAA
CTTACTCTTTCTTTTAAGATAAGTTGTAGACTTCGATGTTTCATGCTCATGTACTTCAAATAATGCATGTTTTAT
AGTTAGTCCCTCATCACTTGAAGTGACTTCTGAGAATTATGCAGAGTCAACATGGATCATTTACAGTGAGATGC
TTTATGGATTGAAGGATATGGTAAAATGTTTATAGTTTACTTTGAAAGTAAAATATACTATGCTTTGGTTTTGAG
GATATTGGATACAAAACCTCTCTTCTTTAGGGCTACTGAGTCTTGATTCTGATCATCAGAAATTTACCAGAAA
CAACTTGCTTCCAATATACCCAATTCTATATGAAGAATTCATGGAGAGTGACTGGCACTGGAAGAGTTTAGTGT
TTCTTGATGCTTGAAAATAAAGTATGTACTGTTTTGAATGIGTTCCAAGTCTCTGCATAAACGATGTATTTG
GGGTCTGGTTGGGCCTGGAAAATGGATGAGCACTTCAGAACAGGTCATTTTCTGATATTGGAAGTGACATGTGG
CCCTATAGGAGGCATGATGTTAGTTAATTACACATTTGCCTACATCTGTGGGAAATGGAGAACAAAGCCATGTGG
GTACTGTAAACACACGTTTATCTTTTGGCCCAATGCCATACATATGGTAGGCATTTAATTACTGATTGTGTTGG
ATAATTTGGGAATTTTCGACTGTGGTAAAATATACATAAAATAACTTATTAATAAAAAAAAAAAAAAAAAA

614/6881
FIGURE 572

MEEYAREPCPWRIVDDCGGAFIMGTIGGGIFQAIKGFRNSPVGVNHLRGSLTAIKTRAPQLGGSFVWGGFLFSM
IDCSMVQVRGKEDPWNSITSGALTGAILAARNGPVAMVGSAAAMGGILLALIEGAGILLTRFASAQFPNGPQFAED
PSQLPSTQLPSSPFGDYRQYQ

615/6881
FIGURE 573

GGCCATGGCGAGCGGCGAGCATTCCCCGGCAGCGGCGGGCCCGGCGCGCTGCACTCCGCGCAGGCTGTGGA
CGTGGCCTCGGCCTCCAACCTCCGGGCCCTTTGAGCTGCTGCACTTGACCTGGACCTGCGGGCTGAGTTCGGGCC
TCCAGGGCCCGGCGCAGGGAGCCGGGGGCTGAGCGGCACCGCGGTCTGGACCTGCGCTGCCTGGAGCCCCGAGGG
CGCCGCGAGCTGCGGCTGGA CTGCAACCGTGCTTGGAGGTGACGGCGGCGGCTGCGGCGGGAGCGGCCCGG
CTCGGAGGAGCCGCTGCGGAGCCCGTGAGCTTCTACACGCAGCCCTTCTCGCACTATGGCCAGGCCCTGTGCGT
GTCCTTCCCGCAGCCCTGCCGCGCCGCCGAGCGCTCCAGGTGCTGCTCACCTACCGCGTCGGGGAGGGACCCGG
GGTTTGCTGGTTGGCTCCCGAGCAGACAGCAGGAAAGAAGAAGCCCTTCGTGTACACCCAGGGCCAGGCTGTCTT
AAACCGGGCCTTCTTCCCTTGCTTCGACACGCCCTGCTGTTAAATACAAGTATTAGCTCTTATTGAGGTCCCAGA
TGGCTTCACAGCTGTGATGAGTGCTAGCACCTGGGAGAAGAGAGGTCCAAATAAGTTCTTCTCCAGATGTGTCA
GCCCCATCCCTCTCTATCTGATAGCTTTGGCCATCGGAGATCTGGTTTCGGCTGAAGTTGGACCCAGGAGCCGGGT
GTGGGCTGAGCCCTGCCTGATTGATGCTGCCAAGGAGGAGTACAACGGGGTGATAGAAGAATTTTTGGCAACAGG
AGAGAAGCTTTTTGGACCTTATGTTTGGGGAAGGTATGACTTGCTCTTCATGCCACCGTCTCTTCCATTGGAGG
AATGGAGAACCCTTGCTGACCTTTGTACCCCCCTGCCTGCTAGCTGGGGACCGCTCCTTGGCAGATGTCATCAT
CCATGAGATCTCCACAGTTGGTTTGGGAACCTGGTCAACCAACGCCAACTGGGGTGAATTCGGCTCAATGAAGG
TTTACCATGTACGCCCAGAGGAGGATCTCCACCATCCTCTTTGGCGCTGCGTACACCTGCTTGGAGGCTGCAAC
GGGGCGGGCTCTGCTGCGTCAGCACATGACATCACTGGAGAGGAAAACCACTCAACAAGCTCCGCGTGAAGAT
TGAACCAGGCGTTGACCCGGACGACACCTATAATGAGACCCCTACGAGAAAGGTTTCTGCTTTGTTTCATACCT
GGCCCACTTGGTGGGTGATCAGGATCAGTTTGACAGTTTTCTCAAGGCCTATGTGCATGAATCAAATTCGGAAG
CATCTTAGCCGATGACTTTCTGGACTTCTACTTGGAATATTTCCCTGAGCTTAAGAAAAAGAGAGTGGATATCAT
TCCAGGTTTTGAGTTTGATCGATGGCTGAATACCCCCGGCTGGCCCCCGTACCTCCCTGATCTCTCCCTGGGGA
CTCACTCATGAAGCCTGCTGAAGAGCTAGCCCCAAGTGTGGGCAGCCGAGGAGCTGGACATGAAGGCCATTGAAGC
CGTGGCCATCTCTCCCTGGAAGACCTACCAGCTGGTCTACTTCTTGATAAGATCCTCCAGAAATCCCCTCTCCC
TCCTGGGAATGTGAAAAAAGTGGAGACACATAACCAAGTATCTCAAATGCCCGGAATGCAGAGCTCCGGCTGCG
ATGGGGCCAAATCGTCCCTTAAGAACGACCACAGGAAGATTTCTGGAAAGTGAAGGAGTTCTTGCATAACCAAGG
GAAGCAGAAGTATACACTTCGCTGTACCACGCAATGATGGGTGGCAGTGAGGTGGCCAGACCCCTCGCCAAGGA
GACTTTTGATCCACCGCTCCAGCTCCACAGCAATGTTGTCAACTATGTCCAGCAGATCGTGGCACCCAAGGG
CAGTTAGAGGCTCGTGTGCATGGCCCTGCCTCTTCAGGCTCTCCAGGCTTTCAGAATAATTGTTTGTTCCTCAA
TTCTGTTCCCTGATCAACTTCTGGAGTTTATATCCCTCAGGATAATCTATTCTCTAGCTTAGGTATCTGTGA
CTCTTGGGCTCTGCTCTGGTGGGAACCTTACTTCTATAGCCCACTGAGCCCGAGACAGAGAACCTGCCACA
GCTCTCCCCGCTACAGGCTGCAGGCACTGCAGGGCAGCGGTATTCTCCTCCCACTAAGTCTCTGGGAAGAAG
TGGAGAGGACTGATGCTCTCTTTTTCTCTTCTGTCTTTTCTTGCTGATTTTATGCAAAGGCTGGCATT
TGATTGTTCTTTTTTTCAGGTTAATCCTTATTTTAATAAAGTTTTCAAGCAAAAATT

616/6881
FIGURE 574

MASGEHSPGSGAARRPLHSAQAVDVASASNFRAFELLHLHLDLRAEFGPPGPGAGSRGLSGTAVLDLRCLEPEGA
AELRLDSHPCLEVTAALRRERPGSEEPPEPVSFYTQPFSSHYQALCVSFPQPCRAAERLQVLLTYRVGEGPGV
CWLAPETAGKKKPFVYTQGGAVLNRAFFPCFDTPAVKYKYSALIEVPDGFATVMSASTWEKRGPNKFFQMCQP
IPSYLIALAIGDLVSAEVGPRSRVWAEPCLIDAAKEEYNGVIEEFLATGEKLFGPYVWGRYDLLFMPPSFPPGGM
ENPCLTFVTPCLLAGDRSLADVIIHEISHSWFGNLVTNANWGEFWLNEGFTMYAQRRISTILFGAAYTCLEAATG
RALLRQHMDITGEENPLNKLRLVKIEPGVDPDDTYNETPYEKGFCFVSYLAHLVGDQDQFDSFLKAYVHEFKFRSI
LADDFLDFYLEYFPELKKKRVDIIPGFEEFDRWLNTPGWPPYLPDLSPGDSLMKPAEELAQLWAAEELDMKAIEAV
AISPWKTYQLVYFLDKILQKSPLPPGNVKKLGDTYPSISNARNAELRLRWGQIVLKNHDEDFWKVKEFLHNQ GK
QKYLPLPYHAMMGSEVAQTLAKETFASTASQLHSNVVNYVQQIVAPKGS

617/6881
FIGURE 575

AGGCTCTATTTAGAGCCGGGTAGGGGAGCGCAGCGGCCAGATACCTCAGCGCTACCTGGCGGAACTGGATTTCTC
TCCCGCCTGCCGGCCTGCCTGCCACAGCCGGACTCCGCCACTCCGGTAGGATTCCCCGCCTGTCAATCCCTAGCC
CAGCTCTTGGGAACTGCAGAGGGGTCCAGAGGATTTCAGTTCTGAACCTGCACACTCCAGTCTAGGATCTCCG
AGCAAGAGCGTAGCCTCATGGCTGCAACCTGTGAGATTAGCAACATTTTATAGCAACTACTTCAGTGCGATGTACA
GCTCGGAGGACTCCACCCTGGCCTCTGTTCCCCCTGCTGCCACCTTTGGGGCCGATGACTTGGTACTGACCCCTGA
GCAACCCCCAGATGTCAATTGGAGGGTACAGAGAAGGCCAGCTGGTTGGGGGAACAGCCCCAGTTCTGGTCGAAGA
CGCAGGTTCTGGACTGGATCAGCTACCAAGTGGAGAAGAACAAGTACGACGCAAGCGCCATTGACTTCTCACCAT
GTGACATGGATGGCGCCACCCTCTGCAATTGTGCCCTTGAGGAGCTGCGTCTGGTCTTTGGGCCTCTGGGGGACC
AACTCCATGCCAGCTGCGAGACCTCACTTCCAGCTCTTCTGATGAGCTCAGTTGGATCATTGAGCTGCTGGAGA
AGGATGGCATGGCCTTCCAGGAGGCCCTAGACCCAGGGCCCTTTGACCAGGGCAGCCCCCTTTGCCAGGAGCTGC
TGGACGACGGTCAGCAAGCCAGCCCCACCACCCCGGCAGCTGTGGCGCAGGAGCCCCCTCCCCGGCAGCTCTG
ACGTCTCCACCGCAGGGACTGGTGCTTCTCGGAGCTCCCACTCCTCAGACTCCGGTGGAAGTGACGTGGACCTGG
ATCCCACTGATGGCAAGCTCTTCCCCAGCGATGGTTTTCTGTGACTGCAAGAAGGGGGATCCCAAGCACGGGAAGC
GGAAACGAGGCCGCCCGGAAAGCTGAGCAAAGAGTACTGGGACTGTCTCGAGGGCAAGAAGAGCAAGCACGGCGC
CCAGAGGCACCCACCTGTGGGAGTTTCATCCGGGACATCCTCATCCACCCGGAGCTCAACGAGGGCCTCATGAAGT
GGGAGAATCGGCATGAAGGCGTCTTCAAGTTCTGCGCTCCGAGGCTGTGGCCCAACTATGGGGCCAAAAGAAAA
AGAACAGCAACATGACCTACGAGAAGCTGAGCCGGGCCATGAGGTACTACTACAAACGGGAGATCCTGGAACGGG
TGGATGGCCGGCGACTCGTCTACAAGTTTGGCAAAAACCTCAAGCGGCTGGAAGGAGGAAGAGGTTCTCCAGAGTC
GGAAGTGAAGGTTGGAAGTATACCCGGGACCAAACTCACGGACCACTCGAGGCCTGCAAACCTTCTGGGAGGAC
AGGCAGGCCAGATGGCCCCCTCCACTGGGGAATGCTCCCAGCTGTGCTGTGGAGAGAAGCTGATGTTTTGGTGTAT
TGTCAGCCATCGTCTGGGACTCGGAGACTATGGCCTCGCTCCCCACCCTCCTCTTGGAATTACAAGCCCTGGG
GTTTGAAGCTGACTTTATAGCTGCAAGTGTATCTCCTTTTATCTGGTGCCTCCTCAAACCCAGTCTCAGACACTA
AATGCAGACAACACCTTCTCCTGCAGACACCTGGACTGAGCCAAGGAGGCCTGGGGAGGCCCTAGGGGAGCACC
GTGATGGAGAGGACAGAGCAGGGGCTCCAGCACCTTCTTTCTGGACTGGCGTTCACCTCCCTGCTCAGTGCTTGG
GCTCCACGGGCAGGGGTCAGAGCACTCCCTAATTTATGTGCTATATAAATATGTCAGATGTACATAGAGATCTAT
TTTTTCTAAAACATTCCCCCTCCCCACTCCTCTCCACAGAGTGCTGGACTGTTCCAGGCCCTCCAGTGGGCTGAT
GCTGGGACCTTAGGATGGGGCTCCCAGCTCCTTTCTCCTGTGAATGGAGGCAGAGACCTCCAATAAAGTGCCTT
CTGGGCTTTTTCTAACCTTTGTCTTAGCTACCTGTGTACTGAAATTTGGGCCTTTGGATCGAATATGGTCAAGAG
GTTGGAGGGGAGGAAAAATGAAGGTCTACCAGGCTGAGGGTGAGGGCAAAGGCTGACGAAGAGGGGAGTTACAGAT
TTCTGTAGCAGGTGTGGGCTTACAGACACATGGACTGGGCTGGGAGGCGAGCAAAGGAAGCAGCTGAGACTGTT
GGAGAACGCTTACAAGACTTCATGCAAGCAAGGACATGAACTCAGAACACTGAGGTGAGAAGCATCCTGCTGTCA
TGACACCGCTCGAGTGACCTTGACCTTGACCAAGTCTGTCTGTTTAGGACTGATTTTTCTATTAGGCTAGGGT
TTGGACCTGATGTTCTCAAGATGTCTAGAATTGCATGGCTGGCCTTGTGGAATAGATGGTTTTGCATTCCAGCCA
AGTGTGCTGTAACTGTATATCTGTAATATGAATCCAGCTTTTGAGTCTGACAAAATCAGAGTTAGGATCTTGT
AAAGGA

618/6881
FIGURE 576

CGGGTACAGGGGGCCCAAGAGCTGGGCTGGCTGTCTCCTGCTCATCCAGCCATGCGGTGGCTGTGGCCCCCTGGCT
GTCTCTCTTGCTGTGATTTTGGCTGTGGGGCTAAGCAGGGTCTCTGGGGGTGCCCCCTGCACCTGGGCAGGCAC
AGAGCCGAGACCCAGGAGCAGCAGAGCCGATCCAAGAGGGGCACCGAGGATGAGGAGGCCAAGGGCGTGACAGAG
TATGTGCCTGAGGAGTGGGCGGAGTACCCCGGCCCATTCACCCTGCTGGCCTGCAGCCAACCAAGCCCTTGGTG
GCCACCAGCCCTAACCCCGACAAGGATGGGGGCACCCAGACAGTGGGCAGGAAGTGGGGGCAATCTGACAGGG
GCACCAGGGCAGAGGCTACAGATCCAGAACCCCTGTATCCGGTGACCGAGAGCTCCTACAGTGCCTATGCCATC
ATGCTTCTGGCGCTGGTGGTGTGTTGCGGTGGGCATTGTGGGCAACCTGTCGGTCATGTGCATCGTGTGGCACAGC
TACTACCTGAAGAGCGCCTGGAACCTCCATCCTTGCCAGCCTGGCCCTCTGGGATTTTCTGGTCCTCTTTTCTGC
CTCCCTATTGTATCTTTCAACGAGATCACCAAGCAGAGGCTACTGGGTGACGTTTCTTGTCGTGCCGTGCCCTTC
ATGGAGGTCTCCTCTCTGGGAGTCACGACTTTTCAGCCTCTGTGCCCTGGGCATTGACCGCTTCCACGTGGCCACC
AGCACCCCTGCCAAGGTGAGGCCCATCGAGCGGTGCCAATCCATCCTGGCCAAGTTGGCTGTATCTGGGTGGGC
TCCATGACGCTGGCTGTGCCTGAGCTCCTGCTGTGGCAGCTGGCACAGGAGCCTGCCCCCACCATGGGCACCCCTG
GACTCATGCATCATGAAACCCCTCAGCCAGCCTGCCCCGAGTCCCTGTATTCACTGGTGATGACCTACCAGAACGCC
CGCATGTGGTGGTACTTTGGCTGCTACTTCTGCCTGCCATCCTCTTCACAGTCACCTGCCAGCTGGTGACATGG
CGGGTGCGAGGCCCTCCAGGGAGGAAGTCAGAGTGCAGGGCCAGCAAGCACGAGCAGTGTGAGAGCCAGCTCAAC
AGCACCGTGGTGGGCCTGACCGTGGTCTACGCCTTCTGCACCTCCCAGAGAACGTCTGCAACATCGTGGTGGCC
TACCTCTCCACCGAGCTGACCCGCCAGACCCTGGACCTCCTGGGCCTCATCAACCAGTTCTCCACCTTCTTCAAG
GGCGCCATCACCCAGTGCTGCTCCTTTGCATCTGCAGGCCGCTGGGCCAGGCCTTCTGGACTGCTGCTGCTGC
TGCTGCTGTGAGGAGTGC GGCGGGGCTTCGGAGGCCTCTGCTGCCAATGGGTCCGACAACAAGCTCAAGACCGAG
GTGTCCTCTTCCATCTACTTCCACAAGCCCAGGGAGTCACCCCACTCCTGCCCCCTGGGCACACCTTGCTTGAGGC
CCCAGTAGGGGTGGGGAGGGAGGGAGAGGCCGCCACCCCGCCGGTGTCTGCTGTTCTTTCCCATAGGTCTTGC
TTTGTGCTGTCTTGCTGTCTAGGGATGGACTTGGTTCTCTTGTCAGGTTTGGGAATCCG

619/6881
FIGURE 577

MRWLWPLAVSLAVILAVGLSRVSGGAPLHLGRHRAETQEQQSRSKRGTEDEEAKGVQQYVPEEWAHEYPRPIHPAG
LQPTKPLVATSPNPKDGGTPDSGQELRGNLTGAPGQRLQIQNPLYPVTESSYSAYAIMLLALVVFAVGIVGNLS
VMCIVWHSYYLKSAWNSILASLALWDFLVLFCLPIVIFNEITKQRLLDVSCRAVFFMEVSSLGVTTFSLCALG
IDRFHVATSTLPKVRPIERCQSILAKLAVIWVGSMTLAVPELLWQLAQEPAPTMGTLDSCIMKPSASLPESLYS
LVMTYQNARMWWYFGCYFCLPILFTVTCQLVTRVRGPPGRKSECRASKHEQCESQLNSTVVGLTVVYAFCTLPE
NVCNIVVAYLSTELTRQTLDLLGLINQFSTFFKGAITPVLLCICRPLGQAFDCCCCCCECGGASEASAANG
SDNKLKTEVSSSIYFHKPRESPLPLPLGTPC

620/6881
FIGURE 578

GCGCGCAGCGCTGGTACCCCGTTGGTCCGCGGTTGCTGCGTTGTGAGGGGTGTCAGCTCAGTGCAATCCCAGGCA
GCTCTTAGTGTGGAGCAGTGAAGTGTGTGGTTTCCTTCTACTTGGGGATCATGCAGAGAGCTTCACGTCTGAAG
AGAGAGCTGCACATGTTAGCCACAGAGCCACCCCAAGGCATCACATGTTGGCAAGATAAAGACCAAATGGATGAC
CTGCGAGCTCAAATATTAGGTGGAGCCAACACACCTTATGAGAAAGGTGTTTTTAAGCTAGAAGTTATCATTCCCT
GAGAGGTACCCATTTGAACCTCCTCAGATCCGATTTCTCACTCCAATTTATCATCCAAACATTGATTCTGCTGGA
AGGATTTGCTGGATGTTCTCAAATTGCCACCAAAAGGTGCTTGGAGACCATCCCTCAACATCGCAACTGTGTTG
ACCTCTATTCAGCTGCTCATGTCAGAACCAACCTGATGACCCGCTCATGGCTGACATATCCTCAGAATTTAAA
TATAATAAGCCAGCCTTCCTCAAGAATGCCAGACAGTGGACAGAGAAGCATGCAAGACAGAAACAAAAGGCTGAT
GAGGAAGAGATGCTTGATAATCTACCAGAGGCTGGTGAAGTCCAGAGTACACAACCTCAACACAGAAAAGGAAGGCC
AGTCAGCTAGTAGGCATAGAAAAGAAATTTTCATCCTGATGTTTAGGGGACTTGTCTGGTTTCATCTTAGTTAATG
TGTCTTTGCCAAGGTGATCTAAGTTGCCTACCTTGAATTTTTTTTAAATATATTTGATGACATAATTTTGTG
TAGTTTATTTATCTTGTACATATGTATTTTGAATCTTTTAAACCTGAAAAATAAATAGTCATTTAATGTTGAAA
AAAAAAAAAAAAAAAAAAAAAAAAAAAA

621/6881
FIGURE 579

MQRASRLKRELHMLATEPPPGITCWQDKDQMDDLRAQILGGANTPYEKGVFKLEVIIPERYPFEPQIRELTPIY
HPNIDSAGRICLDVLKLPKGAWRPSLNIATVLTISIQLLMSEPNPDDPLMADISSEFKYNKPAFLKNARQWTEKH
ARQKQKADEEEMLDNLPEAGDSRVHNSTQKRKASQLVGIEKKFHPDV

622/6881
FIGURE 580A

ATGCTGCAAATGCCGAAGTTAAATGAAATACCTCCGGGGAGGGCAGGCCGAGGGAGGCTCGGGGGAGGGAAGA
TGGCCTGGACAAACAGGTCTGAAGCTGCGAGGCTGGAGTGGAGGGCGAGGGGAGGCGGGCGGCCAGAGCT
CCATGGGACAGCTGGGGAAGCTCCAGGCTACCTACACAACCTGGCCCAGGCTGGTCACGGTGTCCCCCTCCCTG
CTCTGTGCCCTCTCCTTCCAGAAATCCACCATGGAGAGTAAGGATGAGGTCAGCGACACCGACAGTGGCATCATC
CTGCAGTCTGGCCCCGACAGCCCGGTCTCCCCAATGAAGGAGCTGACCCATGCAGTGCACAAGCAGCAGAGGGCC
CTGGAAGCGAGGCTGGAGGCCTGCCCTGGAGGAGCTGAGGAGACTCTGCCTTCGGGAAGCGGAGCTGACGGGCACC
TTGCCAGCGGAGTATCCCCTCAAACCAGGGGAAAAGGCCCCCAAGGTTTCGCCGAGGATCGGAGCGGCTTACAAA
CTGGATGACTGGGCCTTGACAGAGAGGACCCCCCTAAGCAGCCTGGAGCGCCAGCTGGCCCTGCAGCTGCAGATC
ACAGAGGCAGCCCGTTCGGCTGTGCCCTGGAGGAGAACCTCAGCAGGCAGGCTCGGCGGCAGCGGAAGCACTCCATG
CTGCAGGAGGAGAAGAAGCTGCAGGAGCTCCAGCGCTGCCCTGGTCGAGCGGCGGCAATAGCGAGCCACCTCCG
GCTGCTGCTCTCCCCCTGGGCGGAGAGCTCAGTGCCCTCTGATGACAGCTCCCTGTGAGATGGGCTCCTCTGGAG
GAAGAGGAATCCCAAGTGCCAAAACCTCCTCCAGAGTCTCCAGCCCCACCTTCTCGGCCTCTCCCAACCCAAACC
CTTGAGGGTCTGCAGCCAACAGGACCTGAGGCTGGGAGCCGAGAAGGGCTCCAGTCCAGAACAGCCCTGGAAG
GAAACCAGCCTGGACCACCCCTATGAGAAGCCAGGAAGTCTTCTGAGCCCTGGAGCGAGTCCAGCAGCCAGCC
ACCACACCACAGGATGGGCCCAGTGCCCTCCAGCCTGTGGCTTCTGGAGCCTGCCTCCTACCAGTGGTTCCCATC
CGTGGTGTCTTGGCCAGTGGCAGGGCCGACCAAGTGCCTCCAGCCACCCCTGAGATACAGGGGAGGAGGGGCCAG
TCGCAGTCTCTGAGGGTGGATTCTTCCGGGCGGGTCTGAGGGCCGAGGTGCGAGCGCCTTTCGCCGCGCCGC
CCCACTCACTACACGGTGACAGTGCCAGATTCTCTGCTTTCGCCGACCAAGCCCCGCTGCCACGCCGCTGC
CACTCCTGCTCAGAAGACAGTGGCTCTGACGTCTCCAGCATCTCCACCCCACTTCGCCGGGACAGCAGCCCC
GACATCTCCTTTCTGCAGCCTCTCTCCCCTCCCAAGACCATCGTCACCGCGGGCCTGGGTCCCAGCCGCGAGC
AGAGAGCTGGTCGCCCACCACCCCAAGCTACTGCTGCCGCTGGCTATTTCCCGCGGGGCGGTACGTGGTGGTG
GCTGAGAGCCCCCTGCCGCTGGCGAGTGGGAGCTGCGCCGCGCAGCCCCGGGCGCTGCTTACGAGGAGGAGGGC
ACTCCCCTGCGCTACCAGCGTCTGGTGCCCTCCCGCAGCCGCATCGTGCGACGCCCTCCTGAAAGGACAGCCCG
GCAGGCCGGGGCTCAGCAAGGCCCGCGTGTCCGAGGAGCTCAAGTGGTGGACAGCGTGCACGCCCTCCGGAGC
ACCCGCCCCCACTCACTGGACCGCCAAGGAGCTTTCGGGTGAGGAGCCTGCCCTTGGGAGAGAGGGCTTCGGA
CGAGCCCTGGGACCCCGGGCAGAGTGGCCACAGTTTGTGTGCTGCGGAGATCGCTGATGGGGCCCCGTGCA
GTCTTTGTACCTGAAAAAGGAGAGATCATCAGCAGGTTGTAACCTGCGCCCCACGCTGGAAGAACTGTTTCAT
AGAGGGGCTGGGCTGAGACCCCGCCACCCCTGAGTGCCCTCTTTCAGCTCCCCATCCCCATCGCAGGCCGATGAC
CTGGAGCTGAGACCTTTTATTATTTTTTTTTTACACGACTTTTTTCAGAAGCCCTGACCTAAGGATTTATATATG
TGGATTGTCTCAATACCCCTGTGATATGATTATGTTTTATCCCCCAGAGTTTGGCCTACTGGACTTAAGGCCTT
GCCTGTCTGACTGACAGCCTCTATCTCCTTATATAAGACAAGTGGCAGGGGACGAGTGAAGCAGAGTGAAGCCACC
TTGGGAGTTCTCAACGCTCTGTGCTCTGTTCTAAGAAATTCCCTGGGGAAGTGGCCCTGCGCCTCTGTCCCA
CTATTGCTGGAGGCTGGACATGGTACATACTCATGCACATGACTCTCCCCATTTCCAGGTCTCTGGGTACCCC
AGCCTGGGCTGGGGGAGAATCTCTCCCCCTTTCTAATGTGCTCTGTGATGCACACCAAGTGGTAGGTCAA
GGTCAGTATATCCCGGTGGTGTATTGTCTTGCTAGACCCTGCTATTTCTGACCCCTAAATCCTCTTTAGGGA
CCAGTCACTATACCCTGTCTATGCCCTGTGGGCTCCAGACCCCTGAGCTTTGAGTCAGTGGCATCACAGTTG
TAGCCTCAGGGGGTCTGGCTGGGGGCTGGTCCATGCTTGTGGTTAGTGGACAGCAGCCACCCTTTGACAGCTACC
TCTGGGCATCTCAAGGGCTTGACAGCCCCACTGCTCCTTCTAACATTTTGTGTTTTGAGATGGAGTCTCGCTC
TGTGCGCCAGGCTGGAATGCTGTAGCAAGATTTGGGCTCACTGCAACCCCGTCTCCCGGGTTCAAGCGAATCTC
CTGCCTCAGCCTTCCGAGTAGCTGGGATTACAGGCAAGCACCACCATGCCCTGCTAATTTTTTATTTTTAGTAGAG
ATGGGGTTTCATCATGTTGGCCAGGCTGGTCTCGAATCCTGACCTCAAGTGATCCACCTGCCCTTGGCCTTTCAA
AGTGCTGGGATTACAGGCATGAGCCACCGCACCTGGCCCTTCTAACGTTTTTTCATCATAGTCCCAAAAACCAAT
ACTTTACAAGTGGTTTTTGGAAAGGCACCACTTTTGTGGCATGTTCTGGTTGGGAGAGGGAGTCACAGTTCCTACT
CCCCCACCAGCTATGCTTCTGCTCTGAGAAGGTGGTTATTTATACAAACATGGACATACTCACTCCCAAGGGCT
GATGAGATGCTGAATTTTCTTTGGGGGCATTCAATTAATTGTCCAGCTGCAGCGACTGGAGCAAGTCTGGAAGCT
GCCTGTGCTAAGACCACCCAGCTGTCCCTGGGTTCTCATCCTAGGGCCTTCTTGTCTCCAGGTGAGGGGACCTG
CTTCAATGAGAAAGCAACTGAATTGAGGCTAGGAGAGGTAGGGAGAGCTGAGTCTGACTTCACTGTGCAGAAC
TCTCTGCCCCCATGTTACCTGGACTGGAACAGACTGTGAATATAGCAGAAGGTTCGAAGAACTCTGGTGTCTGAC

623/6881

FIGURE 580B

CTAGAAGAGGCACAGTTCTCTCTACTGGAAAGAAAACGATGTAGCCGATTGCACAAGGGTGCCAAGGGAAGACCC
AGGATGGCCCATCAAAGGAACCTGGGGGAGGATGCAGGAGGCTGAAGGGATGCACCTGGCATTCTCTCACTGTG
CTCTTACCGCATCAGCAACCCCCAACTTTTGGGCCTACTCTGCCCCCATGCGTGAATACCCTGCTTGGATGCTG
TGCTTTTCCGGTTTGTCTCTAAGCCCCCTTCTCCAGGGCATGTTGGTTTCCCTGGCCTCTCAGTGTCTTAAGTGG
AGCCCAGAGTGCCTTGTTCTGAGCCAGGAGACGGCTGAGCACTGGCCCTCCACACCTAAGCGTCCTTTACATTAA
CTTATTGGTCTTGATATAACACCTGGTGCCATTGCCAAGTGGCTGTGTCCTCAGCTACAGAGCTGGAATTGTGTGG
GGTTTAGTGCTAAATACTTCAATAAAGTCTGTTTTTTGTGATTGGCTG

624/6881
FIGURE 581

TCCCGGGCCGGGAGCCCAGCAGGCCGGGAGCGGCTGAGGCCACACCCCGGGCCGGGCGCTTCCCTCCGGTGA
ATCATCGCTCGCAGCGGCGGCGCCCGCAGTGGCCGCAGCAGCGCGCGGGCCCTGGCCGCGCCCCAGCCGAGCGC
AGCGCGGAGTCGCCCCGACCTTTCTCTGCGCAGTACGGCCGCGGGACCGCAGCATGGCGGGCATCGCGGCCAAG
CTGGCGAAGGACCGGGAGGCGGCGGAGGGCTGGGCTCCACGACAGGGCCATCAAGTACCTCAACCAGGACTAC
GAGGCGCTGCGGAACGAGTGCCTGGAGGCCGGGACGCTCTTCCAGGACCCGTCCTTCCCGGCCATCCCCTCGGCC
CTGGGCTTCAAGGAGTTGGGGCCCTACTCCAGCAAAACCCGGGGCATCGAGTGGAAAGCGCCCCACGGAGATCTGC
GCTGACCCCCAGTTTATCATTGGAGGAGCCACCCGCACAGACATCTGCCAAGGAGCCCTGGGTGACTGCTGGCTG
CTGGCAGCCATTGCCTCCCTACCTTGAATGAAGAAATCCTGGCTCGAGTCGTCCCCCTAAACCAGAGCTTCCAG
GAAACTATGCAGGGATCTTCACTTCCAGTTCTGGCAATACGGCGAGTGGGTGGAGGTGGTGGTGGATGACAGG
CTGCCACCAAGGACGGGGAGCTGCTCTTTGTGCATTACGCCGAAGGGAGCGAGTTCTGGAGCGCCCTGCTGGAG
AAGGCATACGCCAAGATCAACGGATGCTATGAAGCGCTATCAGGGGGTGCCACCCTGAGGGCTTCGAAGACTTC
ACCGGAGGCATTGCTGAGTGGTATGAGTTGAAGAAGCCCCCTCCCAACCTGTTCAAGATCATCCAGAAAGCTCTG
CAAAAAGGCTCTCTCCTTGGCTGCTCCATCGACATCACCAGCGCCGCGGACTCGGAGGCCATCACGTTTCAGAAG
CTGGTGAAGGGGCACGCGTACTCGGTACCGGAGCCGAGGAGGTTGAAAGTAACGGAAGCCTACAGAACTGATC
CGCATCCGAAATCCCTGGGGAGAAGTGGAGTGGACAGGGCGGTGGAATGACAACTGCCCAAGCTGGAACACTATA
GACCCAGAGGAGAGGGAAAGGCTGACCAGACGGCATGAAGATGGAGAATTCTGGATGTCTTTCAGTGACTTCCTG
AGGCACTATTCCCGCCTGGAGATCTGTAACCTGACCCAGACACTCTCACCAGCGATACCTACAAGAAGTGGAAA
CTCACCAAAATGGATGGGAACTGGAGGCGGGGCTCCACCGCGGGAGGTTGCAGGAACACACATTCTGG
ATGAACCTCAGTACCTGATCAAGCTGGAGGAGGAGGATGAGGACGAGGAGGATGGGGAGAGCGGCTGCACCTTC
CTGGTGGGGCTCATTAGAAGCACCGACGGCGGCAGAGGAAGATGGGCGAGGACATGCACACCATCGGCTTTGGC
ATCTATGAGGTTCCAGAGGAGTTAAGTGGGCAGACCAACATCCACCTCAGCAAAAACCTCTTCTGACGAATCGC
GCCAGGGAGCGCTCAGACACCTTCATCAACCTCCGGGAGGTGCTCAACCGCTTCAAGCTGCCGCCAGGAGAGTAC
ATTCTCGTGCTTCCACCTTCGAACCAACAAGGATGGGGATTCTGTCATCCGGGTCTTTTCTGAAAAGAAAGCT
GACTACCAAGCTGTGATGATGAAATCGAGGCCAATCTTGAAGAGTTCGACATCAGCGAGGATGACATTGATGAT
GGATTACAGGAGACTGTTTGGCCAGTTGGCAGGAGAGGATGCGGAGATCTCTGCCTTTGAGCTGCAGACCATCCTG
AGAAGGGTTCTAGCAAAGCGCCAAGATATCAAGTCAGATGGCTTCAGCATCGAGACATGCAAAATTATGGTTGAC
ATGCTAGATTCCGACGGGAGTGGCAAGCTGGGGCTGAAGGAGTTCTACATTCTCTGGACGAAGATTCAAAAATAC
CAAAAATTACCGAGAAATCGACGTTGACAGGTCTGGTACCATGAATTCCTATGAAATGCGGAAGGCATTAGAA
GAAGCAGGTTTCAAGATGCCCTGTCAACTCCACCAAGTCATCGTTGCTCGGTTTGAGATGACCAGCTCATCATC
GATTTTGATAATTTTGTTCGGTGTTTGGTTTCGGCTGGAACGCTATTCAAGATATTTAAGCAGCTGGATCCCGAG
AATACTGGAACAATAGAGCTCGACCTTATCTCTGGCTCTGTTTCTCAGTACTTTGAAGTTATAACTAATCTGCC
TGAAGACTTCTCATGATGGAATAATCAGCCAAGGACTAAGCTTCCATAGAAATACACTTTGTATCTGGACCTCAAA
ATTATGGGAACATTTACTTAAACGGATGATCATAGCTGAAAATAATGATACTGTCAATTTGAGATAGCAGAAGTT
TCACACATCAAAGTAAAAGATTGTCATATCATTATACTAAATGCAAATGAGTCGCTTAACCCCTTGACAAGGTCAA
AGAAAGCTTTAAATCTGTAAATAGTATACACTTTTACTTTTACACACTTTTCTGTTTCATAGCAATATTAAATCA
GGAAAAAAAATGCAGGGAGGTATTTAACAGCTGAGCAAAAACATTGAGTCACTCTCAAAGGACACGAGGCCCTT
GGCAGGGAATATTTAAAGCAACTTCAAGTTTAAATGACAGCTGTTGATTCTACCAAACAACAGTCCAAGATTACC
ATTTCCCATGAGCCAAGTGGGAAACATGGTATATCATGAAGTAATCTGTCAAGGCATCTGGAGAGTCCAGGAGA
GAAGACTCACCTCTGTGCTTGGGTTAAACAAGAGACAGGTTTTGTAGAATATTGATTGGTAATAGTAAATCGTT
CTCCTTACAATCAAGTTCTTGACCCTATTTCGGCTTATACATCTGGTCTTACAAAGACCAAAGGGATCTGCGCT
TGATCAACTGAACAGTATGCCAAAACAGGCATCCAATTTGTAAACCAATTATGATAAAGGACAAAATAAGCTG
TTTGCCACCTCAAACTTTATGAACCTTACCACCACTAGTGTCTGTCCATGGAGTTAGAGGGGACATCACTTAGA
AGTTCTTATAGAAAGGACACAAGTTTGTTCCTGGCTTTACCTTGGGAAAATGCTAGCAACATTATAGAAATTTT
GCCTTGTGCTTATCTTCTTCCAAATGTACTGTTAAATAAAAATAAAGGTTACCCCATGCAATCACACCATGC
CATGTTTTCTTCTTGGAGGGCAGCCCCACAGGACGGTTTATGAGCACACAATTATAGCTTGTTTTCTACTTTAAC
AAGGTATGCTGCCTCTGTAAATTCATGTATTCAAAGGAAAAGACACCTTGCTTATAATTAAATGTGGAACATA
AAATTTTTTAAATCC

625/6881
FIGURE 582

TCTAATACCTATTGATCTGTTACTTTCTCCCATCACGCTCAGGTGGGAACATCCAGTTGCAGGAAAACAAGCTTA
ACACGCCCCACTGATTCTACATTATGGTGAGTTCCATAATTATTTTATTATATATTACAGTGTAAATGGAATA
AAGTGCCTAATAAATGCAAATGTGCTTACATCTTTTGGCCCAGCTCCTACCTCCCGGCAGCCTCTCCAGGCCCAG
AACTTTCTCCAGTCAGCCTCTACAGACCAAGCTCATGACTCTCAATGGCCTATTTAGGCCCATAACCCTACGTCAC
GGCAGCCTCCGCAGATGAGGCTACTGCCTCACAACAGCCTCCACAGGCACAGCTCCATCGTTACAATGGCCTCTT
TAGACCCAGCTCCTGCCTCCCAGCCTTCTCTCCAGGCCCTGAACTTTCTCAAGTTGACCTCACCAGGCCCAGCTC
ATGCTTCTTTGCAGCCTCTCCAGGCCAGCTCCTGCATCTTGGTGGCCCCCTCCAGGCCAGCCTCTGCCTCCCGT
CAGCCTCTACAGTCCCAACGTCTGCCTCACAGCAGATTCTTCACGCCCAGCTTCTACCTCACTTGGACCCCTCCAG
ACCCAGATGGTGTCTCACTGTGGCATCCTCAGGTGAAGCTCCTGCCTTTCGGCAGCCTCTCCAGGCCAGCTCCT
CCTGCCTCCCAGTGGCCTCTTTCGGCCCAGCCCAGCTCATGCCTCCCGGCGGCCTTCCCAAGCCCCGCTTTTGAC
TTTCGGTGGCCTCTGCAGGCCTCGACAAGGCCAGCCTCCTGCCTCCCGAAGGCCTGCACAGGCCAGCCTCTGC
CTCACAGCGGACTCTC

626/6881
FIGURE 583

GGTATTTTCAGGACAGCCAGGAGGGGGCGCACATCCGCCGAGAACTGTGAGCAAGAGCGTCTGTGCTGAACCATG
GCGCCACCAGAGGGCGCGCGATCCCGCCCCAACCACTTCCCGCTGAGGTGCCAGAAGCAGCGAGGAGCTTCAGC
TTCTCAGGGCAGCAGGAGGGTCTGTAACTTGGTGTCTTCATTGATGATGATTATCCCCACCTTCTAAGAG
ACAAAGACCAACGAGCCACCACAGCCACCAGTCCCAGAACCTGCCAATGCTGGGGAACGGAAATGAGGGAGTTC
AACTCTGGCCCTCACAATCCAGTGGAGGAGACGAACTCATCTGCCTCTGTCCCTCTGGGCACGCCTCATGCCAG
GTGCATCTGTGGACAGGGGCCATGCCCTGGGCTTCCAAAGTTGGAGAGAGCTGCCAGGCTCAGGTCTGAAGGCC
AGAATTCTACAGTAAGTCTACTGAGTCAAGGTGGGAGCAGGGTTCGGTAGCTTCCGAGGCTCTGCGGGAGAATCC
GTTTCTGGCCGTAGAGGTGGCCTGCACTCCGCAGCTTGTGCTGCCCGTCTCGAATGACTGGAGTTTCTGCTTC
TGCTACTACACCTCCCACCTCTCCATCACCTGCTCTGCTCTTACAAGGATCCGAAGAAATGGAATCATCGTATC
GCTGATCTACGTAAACAACTGAAGAATTGTCTGAAAGAAAATATGACATGAACTTATGAATTCAACAGGTGAAG
ATTACAACCTTGATAAATCAACTTTGTCTAGCTCGAGCTGTAAAGCCAAAGGTCCGGTGTATGATCCCATACCTT
TTTCCAGTCTCATGTTGAAGATTTTTATGTAGAAGGCCTTCCCAAAGGAATTTTTTTTTTTTTTTTTTTGA
GATGGAGTTTTCACTCTTATCGCCAGGCTGGGGTGCAATGGCGCAACCTTGCTGGTCACTGCAACCTCTGCCTC
CTGGGTTCAAGAAATCTCCTGCCTTAGCCTCCCAAGTCACTGGGATTACAGGTGCCCACCACCATACCAGGCTA
ATTTTGTATTTTTAGTGGAGATGCGGTTTACCATGTTGGCCGGGCCAGTCTCGAACTCCTGACGTCAAGTGAT
CTTCCCGCCTCGACTCCTGATATCAAGTGATCTTCCCGCCTCGGCCTCCAGAGTGTGAGATTACAGACGTGAA
CCCATGCCTGGCCAGGAATTTTGTTTTTTAGGAAGGCTTCTACTAATGGAATTCCTGGCCTTGAGAGGATGTTA
CTTTAGAAGGAAAGGATTTTTTGTATTAAAGCTGGACCTACCATGAAAGACTTCTGAATCCAGGAAGAGAAA
CTGACTGGGCAACATGTTATTCAGAAACAGGACCTTGCCCTGTCACTCAGGATGGAGTTCAGTGGTCTCATCATG
GCTCATTATAGCCTCAAACCTCCAGGCTCAAGCAATCTACCATGTCTAGCCTTCCAGTAGCTGGGACTACAGAT
ACCGGGTCTCACACTGTCAACCCAGGCTGGAGTACAGTGGCACAACCTCTGCTCACTGCAGCCTCCACCTGCCAGG
CTCAAGCAATCTCCTGCCTCAGCCTCCCGAGTAGCTGGTGATTTCCAGGATGTGAGCTCACAATGACTCAAGCT
GCCACATACTGTTGATTGTGAAATGCCAGTTGAAGCATATGTCTGCGAGCTTAGGGGTGCTACAAGTTGACCAC
TGCAGCAGTAAAGATGACTCTGAAGAATGGCGTGGGTTGGTTCCTTTCAAATGCACTTGAGCAGCGGTCTCCAAC
CACAGGGCCACAGAGCTGGAGCTGGATCTACCATGAAAGACTTGTGAATCCAGGAAGAGAGACTGACTGGGCAAC
ATGTTATTCAGAATCTTCTGTGCCATCCAGGCTGGAGTGCAGTGATGTGATCATAGCTCACTACTTTGGCC
TTCTGAGATCAAGCAATCCTCCCATCTCAACCTCCCAAGTAGCTAGGACTACACAGCATGTCAACCATGCCAG
ATCATTTTGTAGAGTCAGAGTTTACCGTGGTGGCCAGGTTGGCCATGTTGGCCAGATGGGGTCTTCTTTTGT
GCCAGGCTGGCCACAAATCCTGGGCTCAAGTGATCCTCCACCTCGTCCTTGTAGAGATGAGATTAGTTATG
TCGTCCAGGCTGATCTCAAACCTCCTGGGCTAAATCGATTGTCTCACCTCAGCCTCTCAAGTATGTTATGAAGGTT
ATATGTTAGGAAGGGTCCCAGGAGGTAAACCCACACAGATGGGATTTGGGCATAGGTTTGGTTTCCAGGGGGCA
GTGCTGAGCTCTTTGCCAGTGGGAAATGGGATGCTGGTGATTTCCAGTAGGTGACCTCACAGTGACTCAAGCTAC
CACTTACTGTTGATTGTGACGAAATGCCAGCTGAGGCACATGCCTTGGGAGCTAAGTGGTTGCTGCCCTTGACCA
CTGTGAAGACTGGTGTGGGAAGGGTCGTTTTGGATGCACCTGAGCAGGGGTCCCCAACCCCTGAGCCATGGAGCC
GCAAGGAGCCACACAGCAGGAGGTGGGAACATCCAGTTGCAGGAAAACAAGCTTAACACGCCCACTGATTCTACA
TTATGCTCCTACCTCCCGCAGCCTCTCCAGGCCCAGAACTTCTCCAGTCAAGCTCTACAGACCAAGCTCATGA
CTCTCAATGGCCTATTTAGGCCATACCCTACGTACAGGCAGCCTCCGCAGATGAGGCTACTGCCTCACAACAGC
CTCCACAGGCACAGCTCCATCGTTACAATGGCCTCTTTAGACCCAGCTCCTGCCTCCAGCCTTCTCTCCAGGCC
CTGAACCTTCTCAAGTTGACCTCACCAGGCCAGCTCATGCTTCTTTGAGCCTCTCCAGGCCAGCTCCTGCAT
CTTGGTGGCCCTCCAGGCCAGCCTCTGCCTCCCGTCAGCCTCTACAGTCCCAACGTCTGCCTCACAGCAGATT
CTTACAGGCCAGCTTCTACCTCACTGTGGACCCCCAAGCCAAGCTCCCAACCTTTACAGCAGCTTCTACACACCC
AGCTCCTGCCACCCAGTGGCCTCTTTAGGCCAAGCTCATGCTTCACAAGGGCCTTTCCAGGCCCACTTTTGTCT
CATGGCAACCTTCCCTGGCCAGATTCTGCCTGTCTCCAGCAGCCTAGACAGGCCCAGGTCTTGCCTCACACTG
GCCTCTCTACATCCAGCTTATGCCTCACGGTGGCCTCTCCA

627/6881
FIGURE 584

MMIIPHLLRDKDQRATTATSPRTRQCGWTENEGVQLWPSQSSGGDETHLPLSLWARLMPGASVDRGHAPGLPKLE
KAARLRSEGQNSTVSPTESRWEQGR

628/6881

FIGURE 585

GGACGGAAGGAGCGGCGGCGACGGAGGAGGAGGATGGAGGCGGTGGTGTTCGTCTTCTCTCTCCICGATTGTTGC
GCGCTCATCTTCCTCTCGGTCTACTTCATAATTACATTGTCTGATTTAGAATGTGATTACATTAATGCTAGATCA
TGTTGCTCAAAATTAAACAAGTGGGTAAATCCAGAATTGATTGGCCATACCATTGTCAGTGTATTACTGCTCATG
TCATTGCACTGGTTTCATCTTCCTTCTCAACTTACCTGTTGCCACTTGGAATATATATCGATACATTATGGTGCCG
AGTGGTAAACATGGGAGTGTGTTGATCCAACAGAAATACACAATCGAGGGCAGCTGAAGTCACACATGAAAGAAGCC
ATGATCAAGCTTGGTTTCCACTTGCTCTGCTTCTTCATGTATCTTTATAGTATGATCTTAGCTTTGATAAATGAC
TGAAGCTGGAGAAGCCGTGGTTGAAGTCAGCCTACACTACAGTGCACAGTTGAGGAGCCAGAGACTTCTTAAATC
ATCCTTAGAACCGTGACCATAGCAGTATATATTTTCTCTTGGAACAAAAAACTATTTTGCTGTATTTTACCA
TATAAAGTATTTAAAAACAGGAAAAAAAAAAAAAAAAAAAAAAAAAAAAA

629/6881
FIGURE 586

MEAVVFVFSLLDCCALIFLSVYFIITLSDLECDYINARSCCSKLNKWWIPELIGHTIVTVLLMSLHWFIFLLNL
PVATWNIYRYIMVPSGNMGVFDPTTEIHNRGQLKSHMKEAMIKLGFHLLCFFMYLYSMILALIND

630/6881
FIGURE 587

CCCTAATCAAGTACCCCTCTCTCCACCACCTCCTGCCCCACCCCTCCCTGCATCTGGATTCTTTTTGGCATC
CATGTTCAGAAGACAATCGCCCTTTAACTGGACTTGACAGCTGCAATTGCCGGAGCAAACTTAGGAAAGTGTACAG
GATGGAGGATACCTCTTTCCCAAGTGGAGGGAATGCTATTGGTGTGAACTCCGCCCTCATCTAAAACAGATACAGG
CCGTGGAAATGGACCCCTTCTTTAGGGGGTAGTGGTTTTAATGGAAGAAATGAGTGCCCTGCTGGCCAGGAGGAG
AAGAATTGCTGAAAAGGGATCAACAATAGAAACAGAACAAAAAGAGGACAAAGGTGAAGATTACAGAGCCTGTAAC
TTCTAAGGCCCTCTTCAACAAGTACACCTGAACCAACAAGAAAACCTTGGGAAAGAACAAATACAATGAATGGCAG
CAAGTCACCTGTTATCTCCAGACCAAAATCCACACCCCTTATCACAGCCCAAGTGCCAATGGAGTCCAGACGGAAGG
ACTTGACTATGACAGGCTGAAGCAGGACATTTTAGATGAAATGAGAAAAGAATTAACAAAGCTAAAAGAAGAGCT
CATTGATGCAATCAGGCAGGAACCTGAGCAAGTCAAATACTGCATAGAGGAACAGACTAAGGAGAGATAGGACTTT
AATCTGGAGGAAAAATATCCTACAAACAACAACCTGTTCAACACAGCAAAACCCCTACATTTATGAGCTGTAAGAAG
AAAATGGAGACAAACAGAAGGAGGGAACCAACCTACTCTGAAAGCCTTCAGACATTATGACTCTGGTGATAA
GCTCTTTCCCTCTCCGTTTGTCTGCTTTTTTCTGGCCTTTACAACAGAATGGAAGAGAATCATTTAAGAGTTCTCTG
TAACAGTTATGAGAAAATACTAAAACCCATCAGGCAAGATCACCACGCATTGAAATATTTTCATATCAAGATAA
AGTCGCACATTTTCCACAATACATTGCTAAAATAAAGAGGAGAAAGGCTTAGGAAGTTTTTCTGCAGAGAGTGCT
GGTAAAGAATTGAGCAAGTTTGTCTATTGTATTGTAATGTTTCTCTCAGGTTTTGTTCTTCTATCATGTTTGATAT
TCCATGAATAATTGAGATCAGCCCTATGTAAGTTAAGATCATAATATGTGGAACAAATGGAATTGTAAGTGCTTT
CAAAGGGTAATATTTATAAGAAAGTGTCGAAAAATGTTTCTTCAGCTTGAGAAATTTAGAAATGATAGGAAGTT
TCTCGAGTTAGCCTTCATGCAATTTTGTAGATTAAAACATAAAATTTGTCCAGAACTTAAAGATTTAGATGCCTT
CCTAAATTGTTACAATGCTTTACCAAATCTATGACTTCTACATAACACAAACCAGTGGTCAAATGTAAACACTAT
ATTGTAGATTTACTGTAGGTTTTCAACCTTTTTTAGATTTATGCATGTGGACATTTTTATAATGTAATTACAATC
ACCACAAGGTTAGCTTTTTTAATTGCAGACAGTAATGCATGTCACACTAATATGTAGTGGCCTTTTCAAGGCCTA
GTCCAGGGAAAAACATTTTGTAGAGTATAGGGGAGTGGGAGGAAGGGGAGGAATAATTTTTATTTAAAGTTGAT
TTCTGCACTATCTTTTTCTCAGTTACCTGCATGAATAAATAATGAGAAATATTTTGTGACTTTAATTGGTAAATA
TGTTACAAAACCAAGTACTTAATCTTTTACATCATGTCTTCAGCTATTGTATTTTAACCAGTAATTTCAATGGT
CTGAAACATGATTCTGAGCTTCACATAATATCTTAACTGTGGAACCTCAAAGTTTGATCACTGAATTTGGCAGTT
ATTATTACCTAGGTACCCCGCTGTTACACAGGTGTTAGATACGTGTTCTGAATGAAGCTGCTTTTGAATTTT
GTTATGTTGAAATGCAAGAAATAACAATGATGGCAGCAATTAAGGTCACAGAAATCATTAGGTAAAGGAAAACCA
ATGAGGAGTTCTGCAGTTTTCTTTTAATAAGTAAAGTGAGACTTGGGTGGTGGGAAGAAGGAAGGTGGGAAGAAG
GAATTAGACACTCTGCCTGCCACTCTGCGTGTGTGTGCTCTCGCGCACGTGCTGTCTATATGGAAGCCACTCCCT
TTTCTTTCTTTGAAACTGGTAAGGTAAAAATAGGGGAGAAATCCTACATGTTGGAATGATAGCTTTTGGAAAA
TTAAGAACTCTCCAGGCTCTCCATCTTGATTTATGCTTGAGTTGTTATGTGCCATATTGCTTTGAACTCTGA
TTATCAGAAGTTTACTAAAACCTTTGAAATAATTCACCTTCATCTGCTTCTAGATTTTGTACATCTCAGTCCAT
AAAGCAAAGCTTGTGATAGTGTAGTTTTCTAAACGCTGCAATTTGCAGCCTTTACCACTACAAAGAAGTTGG
ATGAGGGATTTTTTTTTCTTTGTCAAATAGTTCTGTTTCTGTAGAAATTTCAATTTTAGATTAACTGTGAT
GGATGGGCTATCATAATTCAAGTATACATTTCTTTTTCTATCAGATATTCATTGTGCATGCAGTAGTAGTAAAAA
CATCAAAGATGCAGCAAGCTTATTAAGTATTATTTCTAAAAGAAATAGGAGGCATTTTCATCTTTATTATTGTA
CTTTTGGTTATGCAACACTTTGATAATATAAACAGTTATGTCCCTATAAATCTGGTCAGCAACCTCTTTTGAT
TTTGTGGGTAAGTTAAATAGTCTGTAGTAGGTAGTACTGGGTACAAGTGGTCCAACTAAGATAAGAGACTA
AAATAAAATGCTAAATCTTAAAAGAACTGGGTTTATGCACTAAACGTTTGTGCCTTGGTCTAATATTAAACATG
ATGATGTGTAAACTGAC

631/6881
FIGURE 588

MSEDNRPLTGAAAIAGAKLRKVSRMEDTSFPSSGNAIGVNSASSKTD TGRGNGPLPLGGSGLMEEMSALLARRR
RIAEKGSTIETE QKEDKGEDSEPVTSKASSTSTPEPTRKPWERTNTMNGSKSPVISRPKSTPLSQPSANGVQTEG
LDYDRLKQDILDEMRKELTKLKEELIDAIRQELSKSNTA

632/6881

FIGURE 589

GGGGCTGCTGGGACTCGCGTCGGTTGGCGACTCCCGGACGTAGGTAGTTTGTGTTGGCCGGGTTCTGAGGCCCTTGC
TTCTCTTTACTTTTTCCACTCTAGGCCACGATGCCGAGTACCAGACCTGGGAGGAGTTCAGCCGCGCTGCCGAGA
AGCTTTACCTCGCTGACCCTATGAAGGCACGTGTGGTTCTCAAATATAGGCATTCTGATGGAACTTGTGTGTTA
AAGTAACAGATGATTTAGTTTGGTGTATAAAACAGACCAAGCTCAAGATGTAAAGAAGATTGAGAAATTC
ACAGTCAACTAATGCGACTTATGGTAGCCAAGGAAGCCCGCAATGTTACCATGGAACTGAGTGAATGGTTTGAA
ATGAAGACTTTGTCTGTACTTAGGAAGTAAATATCTTTGAATTAGAGAAAGTGTGGGACAGAAAGTACTTTA
TGTAAGTAAAGTGGGCTGTTTCAAGCTTAGAGGTCATTTTTGTAAATTTCTTTTAATTACTTTAGAGAGCTAG
GGATGCAAATGTTTTAGTTAGAAAGCCTTTATTTACTTTTGGAAATTGAACAAGAAATGCATCTGTCTTAGAAA
CTGGAGATTATTTGATGTTAGGTAAAACATGTAATTGTTTCTCTGGCAAATTTGTATCAGTAATTTGAAAATGAG
ATATTAGGAAAAACCAATCTTCTTAAATTTAGTTTCATCTTTCTTTAAAGAACATTAAATGTAACCATTTTGTC
AGATCCATGTATTTTGGAGCATAAAATGTATGCTGTTGTGACCAATAAATATAAAATATGGTAATTGGAATTAAC
TCCACACCATAGTATGCATTGTTATACATACTGTGTACCTAATTATGTATAGCAGTGTAGTCTCAATTATATCTG
AAAGTAATTGTGACTAACAAGTATGCTTTGCCTTATTTCCACATTTAACTACCTGTTAATATAAGGGATTTGTA
GTATCAGCTTGTTGAGCAATGACTTTGAATCTAGTTTTCAGTGATCAGAAGCAGCAGTTATTTGAGTGTATGAAT
GGAATGATGATCACTGTGCTATAATGTACTGAAACCACCATATTACAGAAATATTTACTACATATTTTCCATCTG
TAGTTTCTCAGAAGGGCTATGGATTAGTTTGAAGTGTCAAATCCTTGACATCTTCTGTGACACCCCTGCCCATTT
TCTGTCTTTAATTAAACCAAGGTGTTAGGTGTGACTGTCACAAGTGTATGTTTCCAGTAACTAGAAGTACGAT
ATTTGATAATTATATTTGTATTTACCACCTAAATGTAATGTTGATTCCCTCAAGAATGAAATGAAGGCACATACAT
TGAAATATGTTTTGTATAAATTTGTCATGTTGAACAGCATTTTAGCATGGTAAGTTCCTTAGCTATATGAATTT
TGGCATGTTTTCAGAGAGATCAGTAAATAAAATATTAGAT

633/6881
FIGURE 590

MPQYQTWEEFSRAAEKLYLADPMKARVVLKYRHSDGNLCVKVTDDLVCCLVYKTDQAQDVKKIEKFHSQLMRLMVA
KEARNVTMETE

634/6881

FIGURE 591

GCGGCTCCCAGGAGGCGTGAACCGCGGACCATGAGCGTGGGCTTCATCGGGGCCGGCCAGCTGGCCTATGCTCTG
GCGCGGGGCTTCACGGCCGCAGGCATCCTGTGGCTCACAAGATAATAGCCAGCTCCCCAGAAATGAACCTGCCC
ACGGTGTCCGCGCTCAGGAAGATGGGTGTGAACCTGACACGCAGCAACAAGGAGACGGTGAAGCACAGCGACGTC
CTGTTTCTGGCTGTGAAGCCACATATCATCCCCCTTCATCCTGGATGAGATTGGGGCCGACGTGCAAGCCAGACAC
ATCGTGGTCTCCTGTGCGGCTGGTGTACCATCAGCTCTGTGGAGAAGAAGCTGATGGCATTCCAGCCAGCCCC
AAAGTGATTCGCTGCATGACCAACACACCTGTGGTAGTGCAGGAAGGCGCTACAGTGTACGCCACGGGCACCCAT
GCCCTGGTGGAGGATGGGCAGCTCCTGGAGCAGCTCATGAGCAGCGTGGGCTTCTGCACTGAGGTGGAAGAGGAC
CTCATCGATGCCGTCACGGGGCTCAGTGGCAGCGGGCCTGCCTATGCATTCAATGGCTCTGGACGCATTGGCTGAT
GGTGGGGTGAAGATGGGTTTGCCACGGCGCCTGGCAATCCAACTCGGGGCCCAGGCTTGCTGGGAGCTGCCAAG
ATGCTGCTGGACTCGGAGCAGCATCCATGCCAGCTTAAGGACAATGTCTGCTCCCCTGGGGGAGCCACCATCCAC
GCCCTGCACTTTCTAGAGAGTGGGGGCTTCCGCTCTCTGCTCATCAATGCAGTTGAGGCCTCCTGTATCCGAACA
CGAGAGCTACAGTCCATGGCCGACCAAGAAAAGATCTCCCCAGCTGCCCTTAAGAAGACCCTCTTAGACAGAGTG
AAGCTGGAATCCCCACAGTCTCCACACTGACCCCTCCAGCCCAGGGAAGCTCCTCACAAGAAGCCTGGCCCTG
GGAGGCAAGAAGGACTAAGGCAGCATC

635/6881
FIGURE 592

TTTTCTTTGACTTGTTTGTGGATGGAATGTTTACAGACATTTCTAATTACTGCTTTAATTAAATAAATTGGATC
AAAGGCCGTTTCGAGGTATTTTGTGTTTGCCGTTTGTGCTCAGAATTGGCATTGAGAGGTGATTGATACTGCT
AACAAATTTCTAGTACTCTAGTTTGTGTTCAAGAAGAGATTTGGGTAGACGTAATCTTCACCCTTTCAAATTATA
TAACAATACGAACATTATTTTTTATACTGATCATAATTTCCAGATTTGGGGAGGGGGTGATCGTGGCAGGAAAAG
TTGTATGTTTGGTAGTTGCATATGGTGATTTTGTATTTTCAATGCTGGTAGGTAAGTAAGGAGGTCTCTGTACC
ATGGCTCGTACAAAGCAGACTGCCCCGAAATCGACCGGTGGTAAAGCACCCAGGAAGCAACTGGCTACAAAAGCC
GCTCGCAAGAGTGCGCCCTCTACTGGAGGGGTGAAGAACTCATCGTTACAGGCCTGGTACTGTGGCGCTCCGT
GAAATTAGACGTTATCAGAAGTCCACTGAACTTCTGATTGCGAACTTCCCTTCCAGCGTCTGGTGCGAGAAATT
GCTCAGGACTTTAAACAGATCTGCGCTTCCAGAGCGCAGCTATCGGTGCTTTGCAGGAGGCAAGTGAGGCCTAT
CTGGTTGGCCTTTTTGAAGACACCAACCTGTGTGCTATCCATGCCAAACGTGTAACAATTATGCCAAAAGACATC
CAGCTAGCACGCCGCATACGTGGAGAACGTGCTTAAGAATCCACTATGATGGGAAACATTTTCATTCTCAAAAAA
AAAAAAAAAATTTCTCTTCTCTGTTATTGGTAGTTCTGAACGTTAGATATTTTTTTTCCATGGGGTCAAAAGG
TACCTAAGTATATGATTGCGAGTGGAATAAGGGGACAGAAATCAGGTATTGGCAGTTTTTCCATTTTCATTTG
TGTGTGAATTTTAAATATAAATGCGGAGACGTAAAGCATTAATGCAAGTTAAATGTTTCAGTGAACAAGTTTCA
GCGGTTCAACTTTATAATAATTATAAATAAACCTGTAAATTTTCTGGACAATGCCAGCATTGGATTTTTTTA
AAACAAGTAAATTTCTTATTGATGGCAACTAAATGGTGTGTTGTAGCATTTTTATCATACAGTAGATTCCATCCAT
TCACTATACTTTTCTAACTGAGTTGTCCTACATGCAAGTACATGTTTTAATGTTGTCTGTCTTCTGTGCTGTTT
CTGTAAGTTTGCTATTAAAAATACATTAAACTAT

636/6881
FIGURE 593A

AATCTATCAGGGAACGGCGGTGGCCGGTGC GGCGTGTTCGGTGGCGGCTCTGGCCGCTCAGGCGCCTGCGGCTGG
GTGAGCGCACGCGAGGCGGCGAGGCGGCAGCGTGTTTCTAGGTCGTGGCGTCGGGCTCCGGAGCTTTGGCGGCA
GCTAGGGGAGGATGGCGGAGTCTTCGGATAAGCTCTATCGAGTCGAGTACGCCAAGAGCGGGCGCGCTCTTGCA
AGAAATGCAGCGAGAGCATCCCCAAGGACTCGCTCCGGATGGCCATCATGGTGCAGTCGCCCATGTTTGATGGAA
AAGTCCCACACTGGTACCACTTCTCCTGCTTCTGGAAGGTGGGCCACTCCATCCGGCACCCCTGACGTTGAGGTGG
ATGGGTTCTCTGAGCTTCGGTGGGATGACCAGCAGAAAGTCAAGAAGACAGCGGAAGCTGGAGGAGTGACAGGCA
AAGGCCAGGATGGAATTGGTAGCAAGGCAGAGAAGACTCTGGGTGACTTTGCAGCAGAGTATGCCAAGTCCAACA
GAAGTACGTGCAAGGGGTGTATGGAGAAGATAGAAAAGGGCCAGGTGCGCCTGTCCAAGAAGATGGTGGACCCGG
AGAAGCCACAGCTAGGCATGATTGACCGCTGGTACCATCCAGGCTGCTTTGTCAAGAACAGGGAGGAGCTGGGTT
TCCGGCCCCGAGTACAGTGCGAGTCAGCTCAAGGGCTTCAGCCTCCTTGCTACAGAGGATAAAGAAGCCCTGAAGA
AGCAGCTCCCAGGAGTCAAGAGTGAAGGAAAGAGAAAAGGCGATGAGGTGGATGGATGAGTGAAGTGGCGAAGA
AGAAATCTAAAAAGAAAAAGACAAGGATAGTAAGCTTGAAAAAGCCCTAAAGGCTCAGAACGACCTGATCTGGA
ACATCAAGGACGAGCTAAAGAAAGTGTGTTCAACTAATGACCTGAAGGAGCTACTCATCTTCAACAAGCAGCAAG
TGCCTTCTGGGGAGTCGGCGATCTTGGACCGAGTAGCTGATGGCATGGTGTTCGGTGGCCTCCTTCCCTGCGAGG
AATGCTCGGTCAGCTGGTCTTCAAGAGCGATGCCATTACTGCACCTGGGGACGTCACCTGCCTGGACCAAGTGTA
TGGTCAAGACACAGACACCCAACCGGAAGGAGTGGGTAACCCCAAAGGAATTCCGAGAAATCTCTTACCTCAAGA
AATTGAAGGTTAAAAACAGGACCGTATATTCCCCCAGAAACCAGCGCCTCCGTGGCGGCCACGCTCCGCCCT
CCACAGCCTCGGCTCCTGCTGCTGTGAACCTCCTCTGCTTCAGCAGATAAGCCATTATCCAACATGAAGATCCTGA
CTCTCGGGAAGCTGTCCCGGAACAAGGATGAAGTGAAGGCCATGATTGAGAACTCGGGGGGAAGTTGACGGGGA
CGGCCAACAAGGCTTCCCTGTGCATCAGCACCAAAAAGGAGGTGGAAGATGAATAAGAAGATGGAGGAAGTAA
AGGAAGCCAACATCCGAGTTGTGTCTGAGGACTTCTCCAGGACGCTCTCCGCTCCACCAAGAGCCTTCAGGAGT
TGTTCTTAGCGCACATCTTGTCCCTTGGGGGGCAGAGGTGAAGGCAGAGCCTGTTGAAGTTGTGGCCCCAAGAG
GGAAGTCAGGGGCTGCGCTCTCCAAAAAGCAAGGGCCAGGTCAAGGAGGAAGGTATCAACAAATCTGAAAAGA
GAATGAAATTAACCTTAAAGGAGGAGCAGCTGTGGATCCTGATTCTGGACTGGAACACTCTGCGCATGTCTGG
AGAAAGGTGGGAAGGTCTTCAGTGCCACCTTGGCCTGTGGACATCGTTAAAGGAACCACTCCTACTACAAGC
GTAGCAACAACTGGGACGACACAAGGAAAAGGATATTGGATATTAGGTCCTGGGGCCGTGTGGGTACGGTGATCG
CCGGGAACGCTTGGCACTCCAAAAATTTACGAAGTATCCCAAAAAGTTCTACCCCTGGAGATTGACTATGGCC
AGGATGAAGAGGCAGTGAAGAAGCTGACAGTAAATCCTGSCACCAAGTCCAAGCTCCCCAAGCCAGTTCAGGACC
TCATCAAGATGATCTTTGATGTGGAAGTATGAAGAAAGCCATGGTGGAGTATGAGATCGACCTTCAGAAGATGC
CCTTGGGGAAGCTGAGCAAAAGGCAGATCCAGGCCGATACTCCATCCTCAGTGAGGTCCAGCAGGCGGTGTCTC
AGGGCAGCAGCGACTCTCAGATCCTGGATCTCTCAAATCGTTTTACACCCTGATCCCCCAGACTTTGGGATGA
AGAAGCCTCCGCTCCTGAACAATGCAGACAGTGTGCAGGCCAAGGTGGAATGCTTGACAACCTGCTGGACATCG
AGGTGGCCTACAGTCTGCTCAGGGGAGGGTCTGATGATAGCAGCAAGGATCCCATCGATGTCAACTATGAGAAGC
TCAAACTGACATTAAAGGTGGTTGACAGAGATTCTGAAGAAGCCGAGATCATCAGGAAGTATGTTAAGAACACTC
ATGCAACCACACACAATGCGTATGACTTGGAAAGTCATCGATATCTTTAAGATAGAGCGTGAAGGCGAATGCCAGC
GTTACAAGCCCTTTAAGCAGCTTCATAACCGAAGATTGCTGTGGCACGGGTCCAGGACCACCAACTTTGCTGGGA
TCCTGTCCAGGGTCTTCGGATAGCCCCGCTGAAGCGCCCGTGACAGGCTACATGTTTGGTAAAGGGATCTATT
TCGCTGACATGGTCTCCAAGAGTGCCAACCTACTGCCATACGCTCAGGGAGACCCAATAGGCTTAATCCTGTTGG
GAGAAGTTGCCCTTGGAAACATGTATGAACTGAAGCACGCTTCACATATCAGCAAGTTACCCAAGGGCAAGCACA
GTGTCAAAGGTTTGGGCAAACTACCCCTGATCCTTCAGCTAACATTAGTCTGGATGGTGTAGACGTTCTCTTG
GGACCGGGATTTCATCTGGTGTGAATGACACCTCTCTACTATATAACGAGTACATTGTCTATGATATGCTCAGG
TAAATCTGAAGTATCTGCTGAACTGAAATTCATTTTAAAGACCTCCCTGTGGTAATTGGGAGAGGTAGCCGAGT
CACACCCGGTGGCTCTGGTATGAATTCACCGGAAGCGCTTCTGCACCAACTCACCTGGCCGCTAAGTTGCTGATG
GGTAGTACCTGTACTAAACCACCTCAGAAAGGATTTTACAGAAACGTGTTAAAGGTTTCTCTAACTCTCAAGT
CCCTTGTTTTGTGTTGTGTCTGTGGGGAGGGGTTGTTTTGGGGTTGTTTTTGTCTTCTGCCAGGTAGATAAA
ACTGACATAGAGAAAAGGCTGGAGAGAGATTCTGTTGCATAGACTAGTCTATGGAAAAACCAAGCTTCGTTAG
AATGTCGCTTACTGGTTTTCCCGAGGGAAGGAAAAATACACTTCCACCCTTTTTCTAAGTGTTGCTCTTAGT

637/6881

FIGURE 593B

TTTGATTTTGGAAAGATGTTAAGCATTTATTTTGTAGTTAAAAATAAAAACTAATTCATACTATTTAGATTTTCT
TTTTTATCTTGCACTTATTGTCCCTTTTTAGTTTTTTGTTTGCTCTTGTGGTGAGGGGTGTGGGAAGACCA
AAGGAAGGAACGCTAACAATTTCTCATACTTAGAAACAAAAAGAGCTTTCCTTCTCCAGGAATACTGAACATGGG
AGCTCTTGAAATATGTAGTATTAAAAGTTGCATTG

638/6881
FIGURE 594

CGAGCGGGCGGAGCAGGCATTTCCAGCAGTGAGGAGACAGCCAGAAGCAAGCTATTGGAGCTGAAGGAACCTG
AGACAGAAGCTAGTCCCCCTCTGAATTTTACTGATGAAGAACTGAGGCCACAGAGCTAAAGTGACTTTTCCCA
AGGTCGCCCAGCGAGGACGTGGGACTTCTCAGACGTCAGGAGAGTGATGTGAGGGAGCTGTGTGACCATAGAAAG
TGACGTGTTAAAAACCAGCGCTGCCCTCTTTGAAAGCCAGGGAGCATCATTCAATTTAGCCTGCTGAGAAGAAGAA
ACCAAGTGTCGGGATTAGACCTCTCTGCGGCCCAAGTGTTTCGTGGTGCTTCCAGAGGCAGGGCTATGCTCAC
ATTCATGGCCTCTGACAGCGAGGAAGAAGTGTTGATGAGCGGACGTCCCTAATGTGCGCCGAGAGCCCCACGCC
GCGCTCCTGCCAGGAGGGCAGGCAGGGCCCCAGAGGATGGAGAGAACACTGCCCAGTGGAGAAGCCAGGAGAAGCA
GGAGGACGGTGAGGAGGACCCTGACCCTATGTCTGTAGTGGGGTTCCCGGGCGGCCCGCAGGCCTGGAGGAAGA
GCTGACCCTCAAATACGGAGCGAAGCACGTGATCATGCTGTTTGTGCTGTCACTCTGTGCATGATCGTGGTGGT
AGCCACCATCAAGTCTGTGCGCTTCTACACAGAGAAGAATGGACAGCTCATCTACACGACATTCAGTGAGGACAC
ACCTCGGTGGGCCAGCGCCTCCTCAACTCCGTGCTGAACACCCTCATCATGATCAGCGTCATCGTGGTTATGAC
CATCTTCTTGGTGGTGCTCTACAAGTACCGCTGCTACAAGTTTCATCCATGGCTGGTTGATCATGTCTTCACTGAT
GCTGCTGTTCCCTTCACTATATCTACCTTGGGGAGTGCTCAAGACCTACAATGTGGCCATGGACTACCCAC
CCTCTTGCTGACTGTCTGGAACCTTCGGGGCAGTGGGCATGGTGTGCATCCACTGGAAGGGCCCTCTGGTGCTGCA
GCAGGCCTACCTCATCATGATCAGTGCCTCATGGCCCTAGTGTTTCATCAAGTACCTCCAGAGTGGTCCGCGTG
GGTCATCCTGGGCGCCATCTCTGTGTATGATCTCGTGGCTGTGCTGTGTCCCAAAGGGCCTCTGAGAATGCTGGT
AGAACTGCCCAGGAGAGAAATGAGCCCATATTCCTGCCCCTGATATACTCATCTGCCATGGTGTGGACGGTTGG
CATGGCGAAGCTGGACCCCTCCTCTCAGGGTGCCCTCCAGCTCCCTACGACCCGGAGATGGAAGAAGACTCCTA
TGACAGTTTTGGGGAGCCTTCATACCCCGAAGTCTTTGAGCCTCCCTTGAAGTGGCTACCCAGGGGAGGAGCTGGA
GGAAGAGGAGGAAAGGGGCGTGAAGCTTGGCCTCGGGGACTTCATCTTCTACAGTGTGCTGGTGGGCAAGGCGGC
TGCCACGGGCAGCGGGGACTGGAATACCACGCTGGCCTGCTTCGTGGCCATCCTCATTGGCTTGTGTCTGACCCT
CCTGCTGCTTGTGTGTCAAGAAGGCGCTGCCCGCCCTCCCATCTCCATCACGTTTCGGGCTCATCTTTTACTT
CTCCACGGACAACCTGGTGGGCGCTTCATGGACACCCTGGCCTCCCATCAGCTCTACATCTGAGGGACATGGTG
TGCCACAGGCTGCAAGCTGCAGGGAATTTTCATTGGATGCAGTTGTATAGTTTTACACTCTAGTGCCATATATTT
TTAAGACTTTTCTTTCTTAAAAATAAAGTACGTGTTTACTTGGTGAGGAGGAGGCAGAACCAGCTCTTTGGTG
CCAGCTGTTTCATCACCAGACTTTGGCTCCCGCTTTGGGGAGCGCCTCGCTTCACGGACAGGAAGCACAGCAGGT
TTATCCAGATGAAGTGAAGGTGAGATTAGGGCGGGGAGAAGAGCATCCGGCATGAGGGCTGAGATGCGCAAAG
AGTGTGCTCGGGAGTGGCCCTGGCACCTGGGTGCTCTGGCTGGAGAGGAAAAGCCAGTTCCTACGAGGAGTGT
TCCCAATGCTTTGTCCATGATGCTCTTGTATTTTATTGCTTTAGAACTGAGTCCTGTTCTTGTACGGCAGT
CACACTGCTGGGAAGTGGCTTAATAGTAATATCAATAAATAGATGAGTCCTGTTAGAAAAA

639/6881
FIGURE 595

MLTFMASDSEEEVCDERTSLMSAESPTPRSCQEGRQGPEDGENTAQWRSQENEEEDGEEDPDYVCSGVPGRPPGL
EEELTLKYGAKHVIMLFVPVTLCMIVVATIKSVRFYTEKNGQLIYTTFTEDTPSVGQRLNLSVLNTLIMISVIV
VMTIFLVVLYKYRCYKFHGWLIMSSIMLLFLFTYIYLGEVLKTYNVAMDYPTLLLTWNFGAVGMVCIHWKGPL
VLQQAYLIMISALMALVFIKYLPEWSAWVILGAISVYDLVAVLCPKGPLRMLVETAQERNEPIFPALIYSSAMVW
TVGMAKLDPSSQALQLPYDPEMEEDSYDSFGEPSYPEVFEPPLTGYPGEELEEEERGVKLGGLGDFIFYSVLVG
KAAATGSGDWNTTLACFVAILIGLCLTLLLLAVFKKALPALPISITFGLIFYFSTDNLVRPFMDTLASHQLYI

640/6881
FIGURE 596

CGAGCGGCGGCGGAGCAGGCATTTCCAGCAGTGAGGAGACAGCCAGAAGCAAGCTATTGGAGCTGAAGGAACCTG
AGACAGAAGCTAGTCCCCCTCTGAATTTTACTGATGAAGAACTGAGGCCACAGAGCTAAAGTGACTTTTCCCA
AGGTCGCCCAGCGAGGACGTGGGACTTCTCAGACGTCAGGAGAGTGATGTGAGGGAGCTGTGTGACCATAGAAAG
TGACGTGTTAAAAACCAGCGCTGCCCTCTTTGAAAGCCAGGGAGCATATTCATTTAGCCTGCTGAGAAGAAGAA
ACCAAGTGTCCGGGATTACAGACCTCTCTGCGGCCCCAAGTGTTTCGTGGTGCTTCCAGAGGCAGGGCTATGCTCAC
ATTCATGGCCTCTGACAGCGAGGAAGAAGTGTGTGATGAGCGGACGTCCCTAATGTGGCCGAGAGCCCCACGCC
GCGCTCCTGCCAGGAGGGCAGGCAGGGCCCCAGAGGATGGAGAGAACTGCCCAGTGGAAGCCAGGAGAACGA
GGAGGACGGTGAGGAGGACCCTGACCGCTATGTCTGTAGTGGGGTTCCCGGGCGGCCGAGGCCTGGAGGAAGA
GCTGACCCTCAAATACGGAGCGAAGCACGTGATCATGCTGTTTGTGCCTGTCACTCTGTGCATGATCGTGGTGGT
AGCCACCATCAAGTCTGTGCGCTTCTACACAGAGAAGAATGGACAGCTCATCTACAGCATTCACTGAGGACAC
ACCCTCGGTGGGCCAGCGCCTCCTCAACTCCGTGCTGAACACCCTCATCATGATCAGCGTCATCGTGGTTATGAC
CATCTTCTTGGTGGTGTCTACAAGTACCCTGCTACAAGTTCATCCATGGCTGGTTGATCATGTCTTCACTGAT
GCTGTGTTCTCTTACCTATATCTACCTTGGGGAAGTGCTCAAGACCTACAATGTGGCCATGGACTACCCAC
CCTCTTGTGACTGTCTGGAACCTTCGGGGCAGTGGGCATGGTGTGCATCCACTGGAAGGGCCCTCTGGTGTGCA
GCAGGCCTACCTCATCATGATCAGTGCGCTCATGGCCCTAGTGTTCATCAAGTACCTCCCAGAGTGGTCCGCGTG
GGTCATCCTGGGCGCCATCTCTGTGTATGATCTCGTGGCTGTGCTGTGTCCCAAAGGGCCTCTGAGAATGCTGGT
AGAAACTGCCCAGGAGAGAAATGAGCCCATATCCCTGCCCTGATATACTCATCTGCCATGGTGTGGACGGTTGG
CATGGCGAAGCTGGACCCCTCCTCTCAGGGTGCCCTCCAGCTCCCTACGACCCGAGATGGAAGACTCCTATGA
CAGTTTTGGGGAGCCTTCATACCCCGAAGTCTTTGAGCCTCCCTTGACTGGCTACCCAGGGGAGGAGCTGGAGGA
AGAGGAGGAAAGGGGCGTGAAGCTTGGCCTCGGGGACTTCATCTTCTACAGTGTGCTGGTGGGCAAGCGGCTGC
CACGGGCAGCGGGGACTGGAATACCACGCTGGCCTGCTTCGTGGCCATCCTCATTTGGCTTGTGTCTGACCCTCCT
GCTGCTTGTGTGTTCAAGAAGGCGCTGCCCGCCCTCCCCATCTCCATCACGTTTCGGGCTCATCTTTTACTTCTC
CACGGACAACCTGGTGGGCCGTTTCATGGACACCCTGGCCTCCCATCAGCTCTACATCTGAGGGACATGGTGTGC
CACAGGCTGCAAGCTGCAGGGAATTTTCAATTGGATGCAGTTGTATAGTTTTACACTCTAGTGCCATATATTTTA
AGACTTTTCTTTCTTAAAAATAAAGTACGTGTTTACTTGGTGAGGAGGAGGCAGAACCAGCTCTTTGGTGCCA
GCTGTTTCATCACCAGACTTTGGCTCCCGCTTTGGGGAGCGCCTCGCTTCACGGACAGGAAGCACAGCAGGTTTA
TCCAGATGAACTGAGAAGGTCAGATTAGGGCGGGGAGAAGAGCATCCGGCATGAGGGCTGAGATGCGCAAAGAGT
GTGCTCGGGAGTGGCCCCCTGGCACCTGGGTGCTCTGGCTGGAGAGGAAAAGCCAGTTCCCTACGAGGAGTGTTC
CAATGCTTTGTCCATGATGTCTTGTATTTTATTGCCCTTTAGAACTGAGTCCTGTTCTTGTACGGCAGTCA
ACTGCTGGGAAGTGGCTTAATAGTAATATCAATAAATAGATGAGTCCTGTTAGAAAAA

641/6881

FIGURE 597

MLTFMASDSEEEVCDERTSLMSAESPTPRSCQEGRQGPEDGENTAQWRSQENEEDGEEDPDRYVCSGVPGRPPGL
EEELTLKYGAHVIMLFVPVTLCMIVVATIKSVRFYTEKNGQLIYTTFTEDTPSVGQRLNSVLNLTIMISVIV
VMTIFLVVLYKYRCYKFIHGWLIMSSLMLLFLFTYIYLGEVLKTYNVAMDYPTLLLTWNFGAVGMVCIHWKGPL
VLQQAYLIMISALMALVFIKYLPEWSAWVILGAI SVYDLVAVLCPKGPLRMLVETAQERNEPIFPALIYSSAMVW
TVGMAKLDPPSSQALQLPYDPEMEDSYDSFGEPSYPEVFEPPLTGYPGEELEEEEEEERGVKLGLGDFIFYSVLVGK
AAATGSGDWNTTLACFVAILIGLCLTLLLLAVFKKALPALPISITFGLIFYFSTDNLVRPFMDTLASHQLYI

642/6881
FIGURE 598A

TATACTTCGCTACTTGGCTAGAGTTGCAACTACAGCTGGGTTATATGGCTCTAATCTGATGGAACATACTGAGAT
TGATCACTGGTTGGAGTTCAGTGCTACAAAATTATCTTCATGTGATTCCCTTTACTTCTACAATTAATGAACTCAA
TCATTGCCGTCTCTGAGAACATACTTAGTTGGAACTCCTTGAGTTTAGCAGATTATGTGTTTGGGCCACCCT
AAAAGGAAATGCTGCCTGGCAAGAACAGTTGAAACAGAAGAAAGCTCCAGTTCATGTAAAACGTTGGTTTGGCTT
TCTTGAAGCCCAGCAGGCCCTCCAGTCAGTAGGTACCAAGTGGGATGTTTCAACAACCAAAGCTCGAGTGGCACC
TGAGAAAAAGCAAGATGTTGGGAAATTTGTTGAGCTTCCAGGTGCGGAGATGGGAAAGGTTACCGTCAGATTTCC
TCCAGAGGCCAGTGGTTACTTACACATTGGGCATGCAAAAGCTGCTCTTCTGAACCAGCACTACCAGGTTAACTT
TAAAGGGAACTGATCATGAGATTTGATGACACAAATCCTGAAAAAGAAAGGATTTTGAGAAGGTTATCTT
GGAAGATGTTGCAATGTTGCATATCAAACCAGATCAATTTACTTATACTTCGGATCATTTTGAACTATAATGAA
GTATGCAGAGAAGCTAATTCAAGAAGGGAAGGCTTATGTGGATGATACTCCTGCTGAACAGATGAAAGCAGAACG
TGAGCAGAGGATAGAATCTAAACATAGAAAAACCCTATTGAGAAGAATCTACAAATGTGGGAAGAAATGAAAAA
AGGGAGCCAGTTTGGTCAGTCCTGTTGTTTGGCAGCAAAAATTGACATGAGTAGTAACAATGGATGCATGAGAGA
TCCAACCTTTTATCGCTGCAAAATCAACCACATCCAAGAAGTGGAAATAAATACAATGTTTATCCAACATATGA
TTTTGCCTGCCCCATAGTTGACAGCATCGAAGGTGTACACATGCCCTGAGAACAACAGAATACCATGACAGAGA
TGAGCAGTTTTACTGGATTATTGAAGCTTTAGGCATAAGAAAACCATATATTGGGAATATAGTCGGCTAAATCT
CAACAACACAGTGCTATCCAAAAGAAAACCTCACATGGTTTGTCAATGAAGGACTAGTAGATGGATGGGATGACCC
AAGATTTCCCTACGGTTCGTGGTGACTGAGAAGAGGGATGACAGTTGAAGGACTGAAACAGTTTATTGCTGCTCA
GGGCTCCTCACGTTTCAGTCGTGAACATGGAGTGGGACAAAATCTGGGCGTTTAAACAAAAGGTTATTGACCCAGT
GGCTCCACGATATGTTGCATTACTGAAGAAAGAAGTATCCAGTGAATGTACCTGAAGCTCAGGAGGAGATGAA
AGAAGTAGCCAAACACCCAAAGAATCCTGAGGTTGGCTTGAAGCCTGTGTGGTATAGTCCCAAAGTTTTTCATTGA
AGGTGCTGATGCAGAGACTTTTTTCGGAGGGTGAGATGGTTACATTTATAAATTGGGGCAACCTCAACATTACAAA
AATACACAAAAATGCAGATGGAAAAATCATATCTCTTGATGCAAGTTGAATTTGGAAAAACAAAGACTACAAGAA
AACCATAAGGTCATTGGCTTGACAGAGACTACACATGCTCTTCTTATCCAGTAATCTGTGTCACTTATGAGCA
CTTGATCACAAAGCCAGTGCTAGGAAAAGACGAGGACTTTAAGCAGTATGTCAACAAGAACAGTAAGCATGAAGA
GCTAATGTAGGGGATCCCTGCCTTAAGGATTTGAAAAAGGAGATATTATACAACCTCCAGAGAAGAGGATTCTT
CATATGTGATGACACCTTATGAACCTGTAGCCCATATAGTTGCAAGGAAGCCCCGTGTGTTTGTATATACATTCC
TGATGGGCACACAAAGGAAATGCCAACATCAGGGTCAAAGGAAAAGACCAAAGTAGAAGCCACAAAAATGAGAC
CTCTGCTCCTTTTAAAGAAAGACCAACACCTTCTCTGAATAATAATTGTACTACATCTGAGGATTCCTTGGTCCT
TTACAATAGAGTGGCTGTTCAAGGAGATGTGGTTTCGTGAATTAAGCCAAAGAAAGCACCAGGAAGATGTAGA
TGCAGCTGTAAACAGCTTTTGTCTTTGAAAGCTGAATATAAGGAGAAAAGTGGCCAGGAATATAAACCTGGAAA
CCCTCCTGCTGAAATAGGACAGAATATTTCTTCTAATTCCTCAGCAAGTATTCTGGAAAGTAAATCTCTGTATGA
TGAAGTTGCTGCACAAGGGGAGGTGGTTTCGTAAGCTAAAAGCTGAAAAATCCCCCTAAGXXXXXXXXXXXXXXXXX
XX
XX
XXXXXXXXXXXXXXXXXXXXCTTCTCAAGGGGAAGTAGTTCGGGAACTTAAACTGAAAAAGCCCCCTAAGGATCAAGT
AGATATAGCTGTTCAAGAACTCCTTCAGCTAAAGGCACAGTACAAGTCTTTGATAGGAGTAGAGTATAAGCCTGT
GTCGGCCACTGGAGCTGAGGACAAAGATAAGAAGAAGAAAGAAAAAGAAATAAATCTGAAAAGCAGAATAAGCC
TCAGAAACAAAATGATGGCCAAAGGAAAGACCTTCTAAAAACCAAGGAGGTGGGCTCTCATCAAGTGAGCAGG
AGAAGGGCAGGGGCTTAAGAAACAGACCAGGTTGGGTCTTGAGGCAAAAAAGAAAGAAATCTTGCTGATTGGTA
TTCTCAGGTCATCACAAAGTCAGAAATGATTGAATACCATGACATAAGTGGCTGTTATATTCTTCGTCCTGGGC
CTATGCCATTTGGGAAGCCATCAAGGACTTTTTTGATGCTGAGATCAAGAACTTGGTGTGAAAAGTCTACTT
CCCCATGTTTGTGTCTCAAAGTGCATTAGAGAAAGAGAAGACTCATGTTGCTGACTTTGCCCCAGAGGTTGCTTG
GGTTACAAGATCTGGCAAAACCGAGCTGGCAGAACCAATTGCCATTCTGCTACTAGTGAACAGTAATGTATCC
TGATATGCAAAATGGGTACAATCACACAGAGACCTGCCCATCAAGCTCAATCAGTGGTGCAATGTGGTGCCTTG
GGAATTCAGCATCCTCAGCCTTTCTACGTACTCGTGAATTTCTTTGGCAGGAAGGGCACAGTGTCTTGTCTAC
CATGGAAGAGGCAGCGGAAGAGGTCTTGAGATACTTGACTTATATGCTCAGGTATATGAAGAACTCTGGCAAT
TCCTGTTGTTAAAGGAAGAAAGACGGAAGGAAAAATTTGCAGGAGGAGACTATACAACATAAGAGCATT
TATATCTGCTAGTGAAGAGCTATCCAGGGAGGAACATCACATCATTTAGGGCAGAATTTTCCAAAATGTTTGA

643/6881

FIGURE 598B

AATCGTTTTTGAAGATCCAAAGATACCAGGAGAGAAGCAATTTGCCTATCAAACTCCTGGGGCCTGACAACTCG
AACTATTGGTGTTATGACCATGGTTCATGGGGACAACATGGGTTTAGTATTACCACCCCGTGTAGCATGTGTTCA
GGTGGTGATTATTCCTTGTGGCATTACCAATGCACCTTCTGAAGAAGACAAAGAAGCGCTGATTGCAAAATGCAA
TGATTATCGAAGGCGATTACTCAGTGTTAACATCCGCGTTAGAGCTGATTTACGAGATAATTATTCTCCAGGTTG
GAAATTCAATCACTGGGAGCTCAAGGGAGTTCCCATTAGACTTGAAGTTGGGCCACGTGATATGAAGAGCTGTCA
GTTTGTAGCCGTCAGACGAGATACTGGAGAAAAGCTGACAGTTGCTGAAAATGAGGCAGAGACTAAACTTCAAGC
TATTTTGAAGACATCCAGGTCACCCCTTTTCACAAGGGCTTCTGAAGACCTTAAGACTCATATGGTTGTGGCTAA
TACAATGGAAGACTTTCAGAAGATACTAGATTCTGGAAAGATTGTTTCAGATTCCATTCTGTGGGAAATTGACTG
TGAGGACTGGATCAAAAAGACCACTGCCAGGGATCAAGATCTTGAACCTGGTGCTCCATCCATGGGAGCTAAAAG
CCTTTGCATCCCTTCAAACCACTCTGTGAAGTGCAGCCTGGAGCCAAATGTGTCTGTGGCAAGAACCCTGCCAA
GTACTACACCTTATTTGGTTCGAGCTACTGAGGGATGAACGAAAGCCCCCTCTTCAACTCCTCTCACTTTTAAA
GCATTGATATTAGTATCTTCTCAGATACAGACCGTTTATGATTTTTTAAAAAGTAAAAGTTCTAAAATGAAGTC
ACACAGGACAATTATTCTTATGCCTAAGTTAACAGTGGATAAAAGACTTTTCTGTAAACAACTCCAGTAATAAAT
ATCATGAACTA

644/6881
FIGURE 599A

AGTAGCTGCGGCGCAGGGGCGGAGCGAAGGCTGCGGGCGGCGTGGGTACGCGCACACGTTGCATCTTCTTCCTTT
CGCGGGGTCTCCGTAGTTCTGGCACGAGCCAGGCGTACTGACAGGTGGACCAGCGGACTGGTGGAGATGGCGAC
GCTCTCTCTGACCGTGAATTCAGGAGACCCCTCCGCTAGGAGCTTTGCTGGCAGTAGAACACGTGAAAGACGATGT
CAGCATTTCGGTTGAAGAAGGGAAAGAGAATATTCTTCATGTTTCTGAAAATGTGATATTACAGATGTGAATTC
TATACTTCGCTACTTGGCTAGAGTTGCAACTACAGCTGGGTTATATGGCTCTAATCTGATGGAACATACTGAGAT
TGATCACTGGTTGGAGTTCACTGTACAAAATTATCTTCATGTGATTCCCTTACTTCTACAATTAATGAACTCAA
TCATTGCCTGTCTCTGAGAACATACTTAGTTGGAACTCCTTGAGTTTAGCAGATTTATGTGTTTGGGCCACCCCT
AAAAGGAAATGCTGCCTGGCAAGAACAGTTGAAACAGAAGAAAGCTCCAGTTCATGTAAAACGTTGGTTTGGCTT
TCTTGAAGCCCAGCAGGCCTTCCAGTCAGTAGGTACCAAGTGGGATGTTTCAACAACCAAAGCTCGAGTGGCACC
TGAGAAAAAGCAAGATGTTGGGAATTTGTTGAGCTTCCAGGTGCGGAGATGGGAAAGGTTACCGTCAGATTTC
TCCAGAGGCCAGTGGTTACTTACACATTGGGCATGCAAAAGCTGCTCTTCTGAACCAGCACTACCAGGTTAACTT
TAAAGGGAAACTGATCATGAGATTTGATGACACAAATCCTGAAAAAGAAAGGAAGATTTTGAGAAGGTTATCTT
GGAAGATGTTGCAATGTTGCATATCAAACCAGATCAATTTACTTATACTTCGGATCATTGTTGAACTATAATGAA
GTATGCAGAGAAGCTAATTCAGAAGGGAAGGCTTATGTGGATGATACTCCTGCTGAACAGATGAAAGCAGAACG
TGAGCAGAGGATAGACTCTAAACATAGAAAAAACCTATTGAGAAGAATCTACAAATGTGGGAAGAAATGAAAA
AGGGAGCCAGTTTGGTCAGTCCTGTTGTTTGGCAGCAAAATGACATGAGTAGTAACAATGGATGCATGAGAGA
TCCAACCCCTTTATCGCTGCAAAATTCACCACATCCAAGAACTGGAATAAATACAATGTTTATCCAACATATGA
TTTTGCCTGCCCCATAGTTGACAGCATCGAAGGTGTTACACATGCCCTGAGAACAACAGAATACCATGACAGAGA
TGAGCAGTTTTACTGGATTATTGAAGCTTTAGGCATAAGAAAACCATATATTGGGAATATAGTCGGCTAAATCT
CAACAACACAGTGCATCCAAAAGAAAACCTCACATGGTTTGCAATGAAGGACTAGTAGATGGATGGGATGACCC
AAGATTTCTACGGTTTCGTGGTGTACTGAGAAGAGGGATGACAGTTGAAGGACTGAAACAGTTTATTGCTGCTCA
GGGCTCCTCACGTTCACTGCTGAACATGGAGTGGGACAAAATCTGGGCGTTTAAACAAAAGGTTATTGACCCAGT
GGCTCCACGATATGTTGCATTACTGAAGAAAGAAGTGATCCAGTGAATGTACCTGAAGCTCAGGAGGAGATGAA
AGAAGTAGCCAAACACCCAAAGAATCCTGAGGTTGGCTTGAAGCCTGTGTGGTATAGTCCCAAAGTTTTTCATTGA
AGGTGCTGATGCAGAGACTTTTTTCGGAGGGTGAGATGGTTACATTTATAAATTGGGGCAACCTCAACATTACAAA
AATACACAAAATGCAGATGGAAAAATCATATCTCTTGATGCAAAAGTTGAATTTGGAAAACAAAGACTACAAGAA
AACCCTAAGGTCACTTGGCTTGCAGAGACTACACATGCTCTTCTATTCCAGTAATCTGTGTCACTTATGAGCA
CTTGATCACAAAGCCAGTGCTAGGAAAAGACGAGGACTTTAAGCAGTATGTCAACAAGAACAGTAAGCATGAAGA
GCTAATGCTAGGGGATCCCTGCTTAAGGATTTGAAAAAAGGAGATATTATCAACTCCAGAGAAGAGGATTCTT
CATATGTGATCAACCTTATGAACCTGTTAGCCCATATAAGTTGCAAGGAAGCCCCGTGTGTTTTGATATACATTCC
TGATGGGCACACAAAGGAAATGCCAACATCAGGGTCAAAGGAAAAGACCAAAGTAGAAGCCACAAAAATGAGAC
CTCTGCTCCTTTTAAAGAAAGACCAACACCTTCTCTGAATAATAATTGTACTACATCTGAGGATTCTTGGTCCT
TTACAATAGATGGCTGTTCAAGGAGATGTGGTTTCGTGAATTAAAAGCCAAGAAAGCACCAAGGAAGATGTAGA
TGCAGCTGTAAAACAGCTTTTGTCTTTGAAAGCTGAATATAAGGAGAAAACCTGGCCAGGAATATAAACCTGAAA
CCCTCCTGCTGAAATAGGACAGAATATTTCTTCTAATTCCTCAGCAAGTATTCTGGAAAGTAAATCTCTGTATGA
TGAAGTTGCTGCACAAGGGGAGGTGGTTCTGAAGCTAAAAGCTGAAAAATCCCCTAAGGCTAAAATAAATGAAGC
TGTAAGATGCTTACTGTCCCTGAAGGCTCAGTATAAAGAAAAAACTGGGAAGGAGTACATACCTGGTCAGCCCCC
ATTATCTCAAAGTTTCGGATTCAAGCCCAACCAGAAATTCCTGAACCTGCTGGTTTAGAAAACACCAGAAGCGAAAGT
ACTTTTTGACAAAGTAGCTTCTCAAGGGGAAGTAGTTCGGAAACTTAAACTGAAAAAGCCCCTAAGGATCAAGT
AGATATAGCTGTTCAAGAACTCCTTCAGCTAAAGGCACAGTACAAGTCTTTGATAGGAGTAGAGTATAAGCCTGT
GTCGGCCACTGGAGCTGAGGACAAAGATAAGAAGAAGAAAGAAAAAGAAATTAATCTGAAAAGCAGAATAAGCC
TCAGAAACAAATGATGGCCAAAGGAAAGACCCCTTCTAAAAACCAAGGAGGTGGGCTCTCATCAAGTGGAGCAGG
AGAAGGGCAGGGGCCTAAGAAACAGACCAGGTTGGGTCTTGAGGCAAAAAAAGAAAGAAATCTTGCTGATTGGTA
TTCTCAGGTCACTACAAAGTCAGAAATGATTGAATACCATGACATAAGTGGCTGTTATATTCTTCGTCCTGGGC
CTATGCCATTTGGGAAGCCATCAAGGACTTTTTTGATGCTGAGATCAAGAAACTTGGTGTTGAAAACGCTACTT
CCCCATGTTTGTGTCTCAAAGTGCATTAGAGAAAAGAAAGACTCATGTTGCTGACTTTGCCCCAGAGGTTGCTTG
GGTTACAAGATCTGGCAAAACCGAGCTGGCAGAACCAATTGCCATTGCTCCTACTAGTGAACAGTAATGTATCC
TGCATATGCAAAATGGGTACAGTCACACAGAGACCTGCCCATCAAGCTCAATCAGTGGTGCAATGTGGTGCGTTG

645/6881

FIGURE 599B

GGAATTCAAGCATCCTCAGCCTTTCCTACGTACTCGTGAATTTCTTTGGCAGGAAGGGCACAGTGCTTTTGCTAC
CATGGAAGAGGCAGCGGAAGAGGTCTTGAGATACTTGACTTATATGCTCAGGTATATGAAGAACTCCTGGCAAT
TCCTGTTGTTAAAGGAAGAAAGACGGAAAAGGAAAAATTTGCAGGAGGAGACTATACAACTACAATAGAAGCATT
TATATCTGCTAGTGGAAGAGCTATCCAGGGAGGAACATCACATCATTTAGGGCAGAATTTTCCAAAATGTTTGA
AATCGTTTTTTGAAGATCCAAAGATACCAGGAGAGAAGCAATTTGCCTATCAAACTCCTGGGGCCTGACAACTCG
AACTATTGGTGTATGACCATGGTTCATGGGGACAACATGGGTTTAGTATTACCACCCCGTGTAGCATGTGTTCA
GGTGGTGATTATTCCTTGTGGCATTACCAATGCACTTCTGAAGAAGACAAAGAAGCGCTGATTGCAAAATGCAA
TGATTATCGAAGGCGATTACTCAGTGTAAACATCCGCGTTAGAGCTGATTACGAGATAATTATTCTCCAGGTTG
GAAATTCAATCACTGGGAGCTCAAGGGAGTTCCCATTAGACTTGAAGTTGGGCCACGTGATATGAAGAGCTGTCA
GTTTGTAGCCGTCAGACGAGATACTGGAGAAAAGCTGACAGTTGCTGAAAATGAGGCAGAGACTAACTTCAAGC
TATTTTGAAGACATCCAGGTCACCCCTTTTCAAGGGCTTCTGAAGACCTTAAGACTCATATGGTTGTGGCTAA
TACAATGGAAGACTTTCAGAAGATACTAGATTCTGGAAAGATTGTTTCAGATTCCATTCTGTGGGAAATTGACTG
TGAGGACTGGATCAAAAAGACCACTGCCAGGGATCAAGATCTTGAACCTGGTGCTCCATCCATGGGAGCTAAAAG
CCTTTGCATCCCCCTTCAAACCACTCTGTGAAGTGCAGCCTGGAGCCAAATGTGTCTGTGGCAAGAACCCTGCCAA
GTACTACACCTTATTTGGTTCGAGCTACTGAGGGATGAACGAAAGCCCCCTCTCAACTCCTCTCACTTTTTAAA
GCATTGATATTAGTATCTTCTCAGATACAGACCGTTTTATGATTTTTTAAAAAGTAAAGTTCTAAAATGAAGTC
ACACAGGACAATTATCTTATGCCTAAGTTAACAGTGGATAAAAGACTTTTCTGTAAACAACCTCCAGTAATAAAT
ATCATGAACATAATATGGTTTAAAAAAAAAAAAAAAAAAAAAGG

646/6881
FIGURE 600

CCCAGGATGTAGAGCTGGCAGTGCCTGACGGCGCGTCTGACGCGGAGTTGGGTGGGGTAGAGAGTAGGGGGCGGT
AGTCGGGGGTGGTGGGAGAAGGAGGAGGCGGCAAAATCACTTATAAATGGCGCGGAAGCAGGACCCGAAGCCTAAA
TTCCAGGAGGGGTGAGCGAGTGCTGTGCTTTTCATGGGCCTCTTCTTTATGAAGCAAAGTGTGTAAAGGTTGCCATA
AAGGACAAACAAGTGAAATACTTCATACATTACAGTGGTTGGAATAAAAATTGGGATGAGTGGGTTCCGGAGAGC
AGAGTACTCAAATACGTGGACACCAATTTGCAGAAACAGCGAGAACTTCAAAAAGCCAATCAGGAGCAGTATGCA
GAGGGGAAGATGAGAGGGGCTGCCCCAGGAAAGAAGACATCTGGTCTGCAACAGAAAAATGTTGAAGTGAAAACG
AAAAAGAACAACAGAAAAACCTTGAAATGGAGATGGTGGCAGTACCAGTGAGACCCCTCAGCCTCCTTGGAAG
AAAAGGGCCCCGGGTAGATCCTACTGTTGAAAATGAGGAAACATTTCATGAACAGAGTTGAAGTTAAAGTAAAGATT
CCTGAATAGCTAAAACCGTGGCTTGTTGATGACTGGGACTTAATTACCAGGCAAAAACAGCTCTTTTATCTTCCT
GCCAAGAAGAATGTGGATTCCATTCTTGAGGATTATGCAAATTACAAGAAATCTCGTGGAACACAGATAATAAG
GAGTATGCGGTAAATGAAGTTGTGGCAGGGATAAAAGAATACCTCAACGTAATGTTGGGTACCCAGCTACTCTAT
AAATTTGAGAGACCACAGTATGCCGAAATTCCTGCAGATCATCCCGATGCACCCATGTCCCAGGTGTATGGAGCG
CCACATCTCCTGAGATTATTTGTACGAATTGGAGCAATGTTGGCCTATACACCTCTGGATGAGAAGAGCCTTGCT
TTATTACTCAATTATCTTCACGATTTCTTAAAGTACCTGGCAAAGAATTCTGCAACTTTGTTTAGTGCCAGCGAT
TATGAAGTGGCTCCTCCTGAGTACCATCGGAAAGCTGTGTGAGAGGCACTCTCACTCACTTATGTTTGGATCTCC
GTAAACACATTTTGTTCCTAGTCTATCTCTGTACAAACAATGTGCTTTGAAGATGTTAGTGATAACAATTGA
TGTTTGTTTCTGTTTGATTTTAAACAGAGAAAAATAAAGGGGTAATAGCTCCTTTTTTCTTTTCTTTTTTTTT
TCATTTCAAAGTTGCTGCCAGTGTTCATGATGGACAACAGAGGGATATGCTGTAGAGTGTATTATGCCTAG
TTGACAAAGCTGCTTTTGAATGCTGGTGGTTCTATTCTTTGACACTATGCACTTTTATAATACATGTTAATGCT
ATATGACAAAATGCTCTGATTCCTAGTGCCAAAGGTTCAATTCAGTGTATATAACTGAACACACTCATCCATTG
TGCTTTTTTTTTTTTTTTTATGGTGCTTAAAGTAAAGAGCCCATCCTTTGCAAGGCATCCATGTTGTTACTTAGG
CATTTTATCTTGGCTCAAATTGTTGAAGAATGGTGGCTTGTTCATGGTTTTTGTATTTGIGTCTAATGCACGTT
TTAACATGATAGATGCAATGCACTGTGTAGCTAGTTTTCTGGAAAAGTCAATCTTTTAGGAATTGTTTTTCAGAT
CTTCAATAAAATTTTTCTTTAAATTC

647/6881

FIGURE 601

GAGGAGGCGGCAAATCACTTATAAATGGCGCGGAAGCAGGACCCGAAGCCTAAATTCCAGGAGGGTGAGCGAGTG
CTGTGCTTTTCATGGGCCTCTTCTTTATGAAGCAAAGTGTGTAAAGGTTGCCATAAAGGACAAACAAGTGAAATAC
TTCATACATTACAGTGGTTGGAATAAAAATTGGGATGAGTGGGTTCCGGAGAGCAGAGTACTCAAATACGTGGAC
ACCAATTTGCAGAAACAGCGAGAACTTCAAAAAGCCAATCAGGAGCAGTATGCAGAGGGGAAGATGAGAGGGGCT
GCCCCAGGAAAGAAGACATCTGGTCTGCAACAGAAAAATGTTGAAGTGAAAACGAAAAAGAACAAACAGAAAACA
CCTGGAAATGGAGATGGTGGCAGTACCAGTGAGACCCCTCAGCCTCCTTGGAAGAAAAGGGCCCGGGTAGATCCT
ACTGTTGAAAATGAGGAAACATTCATGAACAGAGTTGAAGTTAAAGTAAAGATTCCTGAATAGCTAAAACCGTGG
CTTGTTGATGACTGGGACTTAATTACCAGGCATTATGCAAATTACAAGAAATCTCGTGGAACACAGATAATAAG
G

648/6881
FIGURE 602

AATCACTTATAAATGGCGCGGAAGCAGGACCCGAAGCCTAAATTCCAGGAGGGTGAGCGAGTGCTGTGCTTTCAT
GGGCCTCTTCTTTATGAAGCAAAGTGTGTAAAGGTTGCCATAAAGGACAAACAAGTGAATACTTCATACATTAC
AGTGGTTGGAATAAAAAATTGGGATGAGTGGGTTCCGGAGAGCAGAGTACTCAAATACGTGGACACCAATTTGCAG
AAACAGCGAGAACTTCAAAAAGCCAATCAGGAGCAGTATGCAGAGGGGAAGATGAGAGGGGCTGCCCCAGGAAAG
AAGACATCTGGTCTGCAACAGAAAAATGTTGAAGTTACGAATTGGAGCAATGTTGGCCTATACACCTCTGGATGA
GAAGAGCCTTGCTTTATTACTCAATTATCTTCACGATTTCTAAAGTACCTGGCAAAGAATTCTGCAACTTTGTT
TAGTGCCAGCGATTATGAAGTGGCTCCTCCTGAGTACCATCGGAAAGCTGTGTGAGAGGCACTCTCACTCACTTA
TGTTTGGATCTCCGTAAACACATTTTGTTCCTTAGTCTATCTCTTGTACAAACAATGTGCTTTGAAGATGTTAGT
GTATAACAATTGATGTTTGTTCCTGTTTGATTTTAAACAGAGAAAAATAAAAGGGTAATAGCTCCTTTTTTCTT

649/6881
FIGURE 603

TTTTGCGCTCGGACCTTCGCCAGAGGGGCCGGGACATCATGACGGTGGGAGCCAGGCTCCGAAGCAAGCGGAGA
GCAGCCTCCTGCGCCGCGGGCCCCGAGGGCGAGGGCGAACCAGGGGGACGAGGAGGCGGCCATCCTGGAGC
ACCTGGAGTACGCGGACGAGGCGGAGGCGGCGCCGAGAGCGGGACGAGCGCGGGACGAGCGGGCCCCGGGA
CCCCGGGCGCGCGGAGGGTGCACTTCGCCCTCCTGCCCGAGCGCTACGAGCCACTGGAGGAGCCGGCGCCGAGCG
AGCAGCCCAGGAAGAGGTACCGGAGGAAGCTGAAGAAGTACGGCAAGAATGTCGGGAAGGTCATCATCAAAGGAT
GCCGCTACGTGGTCATCGGCCCTGCAAGGCTTCGCTGCAGCCTACTCCGCCCCGTTTGCGGTAGCCACCAGCGTGG
TATCCTTCGTGCGCTAAATGGGAGCTGCTGTGGCAGGTGCCCCAGAGTGAACGGGAGCCCCTGCTGTGGGAACCTT
TGTGAATCCTGGAGCATCTCAGACTTGAACACACAGCATATTTGGAAGAGAAAACATGCCTTTCTTTGTTGAATC
ACATTAGTATGATGAGTGAGTCATCCCTGCCATCTGCTGAGCTTCTCACATCTCTCAGTCACACGTGGACCCAG
TGGTCAATCCTGCAGAGAATTCGGCGGAGGTTAGGTTTGGGAGTGGAGCTAGCGTGCTAAAGCCAGAGCCTTCAC
GTGAAGGTGGCAGGCACTGGGGCGGAAGCCAACACTCAACAGATGCAAGCAGTGTGGGTGTGCAGCAGAACAGTG
ATCTTGGGGGAGGAAGAGGATGTTACTAGAGTCAGATGATTTGCTGTATTCTCCTGAAAGGTCGTAGGCTGACAG
GCGCTCACATTCCTTGGCTGCCCTCGGTTCTGAGGGCAGCTAAGGAGCTGTTTATTCTCAAGTCATGCTCCCCGA
TCTCCTTCCTCTACCACTCTGTCAACAGGAGTTAATTACAGGCTTGAGGAGAAGAAAGGAAGATATCTT
GATGCTTTGAAAACCTGTGTTGGCAGTGTGGCATGACTGTTTAAAGTAGATAAAACCTTGTCATTTTACCCCATCC
CTGCATGACTGTGAAGCTGGCGAGGAAGGAGGAAGAAGGGCAAGTTCAGATGCAGGCTGGGTGGCTGGGACAGGT
TGGCTAAGGGACTACTCTGGAGGGCTCTTCTGCCTGGCATTGCCCACTTCGGCCAGCCACGTGTTTGCAGCGAC
CAGAGTCCCTGC AAAAGGTGTGGCTGGCTGTGGTCAGGGTGCTACTAGCACCATCAGCGCACTCCCGCCATTGGCT
CAGCTCCTCTCTGCCAGTCCAACCTAAGAGTGCTTTGTCTTGGGTGGACATAGGGGCTGAGAGAGATGGGGGAG
ACATAACACCCAGGAATGAAAATACAGATTTAGAGAAGGAACCAGTAAGTAGGAGACAGATGTGAAGGAAATGGA
AATGAGGCAAGAGGACATTGGAAGAGAGAAGTTTGTCTGCCAGGAGCCAGGTCTGGAGCATCAGTGTGAGGGAGT
TCAGGTAGGCTGGGCCTGTGCCCTTAGGTAGGGACAAGGGAGGCTGGGTAGCCAGGGCTGGTGCTTAAACCCCT
GAGGCCATGAGCTCATTGGCTGCCCTTGTAGCATCCTGTCTTCTTCTGTGCTGCCTGGTTTGATCTCATCTCACC
TGGATTCAAAGGGTAAGGTGGGCATGGGTCTTGGGCCTGACACCCACCAAGGATGACCTGTGGACTGCCATCGGA
TGCTGAACAGGGAGATGAAAGGAGGTCTCTTACCATACCCCCTCTGCCAACCCCCCAGTAGGCCACTGTTCTGAC
TTTGTTCAGAAATATCCAGAAATCAAAGGGGCTGTGCTGAACAGTCTGCAGGACCAGTGACAGCACCTACCT
GTTGTCCCAAGGCATACAAAGGAGGCCTCAACGCTCATGCTTCTCTAATCAAGCCCTACCAAGACAGACAGAAAG
ACAGACAGAAAAAAGGAAGGGGTAGAGGAGAAGGTTGAAGCTGTGGAGCTAGACTCTGCTTCACTTCCTGAAGCT
TCAACTTCATGTGGAAGATTCACTGGGACCCAATTCTGCAATTGTTAATATTTGTGAGGAAAAGTGAACAAAGTG
ATCTGGTTTTAGCCCAGATGATGAAAGTGATATGGCACATTTTACACACGTGAGATAATTACAGCTTGCCCCA
CAACACTGGGTGTTGGAGAAAGGGAGAGATAGTCATAAGTGGAAGAAAAAGCCAAGCATAGTGAGTGGGAAGGAG
AGTGAGAGCCTGTGCAGGCTGCTGACGAGCCCCAGGCAGCCACAAGTTTCTCGTGGGGAGATGGAGGCAGAGCC
CAGGGTAGGGGACAGAGCTGCTGGGGCCTTTCTTGGCTGGGAATCTGTCCAGGAAGAGCTTCCCCACTCCCAT
CCCCAAATTGGA AAAACCGTACATTCAAGCCTGTTTGGCCCTGAAATCTTAAGAATCTGGTTAAGAATTAAC
CACTAATGTCAAAGTCAAACCTCCTAGGGGTTGTCTGGGAGTCAGGTTACGGGTACAGAAGATGAATCTCA
GATGTCACTCAACCTGAGCCGTCATTCTGTGTGGCAGGGCTGCCCTGGGTTTCTCTTACTCAATCCCTGGAGTGT
AAGCATTTGGATTGTGTACAGATTACCTTTTTACCTTTTCTTTCTTTTTTTCTTTTTTTCAATATCAGTGCC
CACACCTTACTGAGTATTGAGTTTTAGAGCTTTGCTTGATGTGCTTGACCAAGAGACTTCTTTTGTATCCTTTT
CTTGTCTATGATGTAAATAAAAGCCTCGATTTATGT

650/6881
FIGURE 604

MTVGARLRSKAESSLLRRGPRGRGRTEGDEEAAAILEHLEYADEAEAAAESGTSAADERGPGTRGARRVHFALLP
ERYEPLLEPPAPSEQPRKRYRRKLKKYGKNVGKVIKGCYVVGIGLQGFAAAYSAPFAVATSVVSFVR

651/6881
FIGURE 605

TCTAATACCTATTGATCTGTCACCTTTCTCCCATCACGCTCAGGTGGGACCATCCAGTTGCAGGAAAACAAGCTTA
ACATGCCCACCTGATTCTACATTATGGTGAGTTCTATAATTATTTATTATATATTACAGTGTAATAATGGAAATA
AAGCACCTAATAAATGTAAATGTGCTTAAATCTTTTGTCCCAGCTCCTACCTCCCGGCAGCCTCTCCAGGCCCAG
AACTTTCTCCAGTCAGCCTCTACAGACCAAGCTCATGACTCACAATGGCCTATTTAGGCCCATACCCTACCTCAC
GGCAGTCTCCGCAGATGAGCCTACTGCCTCACACAGCCTCCACAGGCACAGCTCCATCGTTACAATGGCCTCTT
TAGACCCAGCTCCTGCCTCCCAGCCTTCTCTCCAGGCCCTGAACTTTCTCAAGTCGACCTCACCAGGCCAGTTC
ATGCTTCTTTGCAGCCTCTCCAGGCCAGCTCCTGCATCTTGGTGGCCCCCTCCAGGCCAGCCTCTGCCTCCCGT
CGGCCTCTACAGTCCCAACATCTGCCTCACAGCAGATTCTTCAGGCCAGCATCTGCCTCACTGTGGACCCCCCA
GACCCAGATGGTGTCTCACTGTGGCATCCTCAGGTGAAGCTCCTGCCTTTCTGCAGCCTCTCCAGGCCAGCTCC
TCCTGCCTCCAGTGCCCTCTTTCAGCCCAGCCAGCTCATGCCTCCTGGCGGCCTTCCCAGGCCCTGCTTTTGA
CTTTTGGTGGCCTCTGCAGGCCTCGACAAGGCCCGGCCTCCTGCCTCCCGAA

652/6881
FIGURE 606

GGCACGAGGGCTCTGTGGAGCTGCGCCTGCCAACTCATTGCTTTCTGCTGTGCTGTGATCCTGCAGTCAGTTGCC
GCTGTTGCTACTGCAATCATTTCACCAAACCTGAGCCAGAGAAGTTCCCTCTTTTAACATAAAGAGTCAAGAAAT
TGGAGCATGGCTATGAGCAATGGAAACAATGATTTTGTGGTTCTGAGCAACAGCAGCATCGCAACCAAGTGTGCT
AACCCGAGTCCCCTCACCCCTGTGATGGAGACCATGCAGCCAGCAGCTCACACCCAAAGAAGCAACAAGAACA
AAAGTGAGTCCAAATGGATGCCTGCAACTTAATGGCACGGTCAAATCATCCTTTCTGCCTTTAGACAACCAAAGA
ATGCCTCAGATGTTACCCCAATGCTGCCATCCTTGCCCATACCATCACCCCTTGACTAGCCATAGCAGTCACCAA
GAGTGCCATCCCGAGGCTGGCCCTGCAGCACCCCTCTGCTTTGGCCTCGTGTTGCATGCAGCCACACTCCGAGTAT
TCTGCATCTCTTTGTCCAAATCATTACCTGTGTATCAGACTACGTGCTGTCTTCAGCCCTCTCCATCCTTCTGC
CTGCATCATCCGTGGCCTGACCATTTTCAGCATCAGCCTGTGCAACAGCACATAGCCAACATAAGACCATCCAGA
CCTTTCAAGTTGCCAAAAGTTATGCAGCCCTGATAGCCGACTGGCCGGTGGTGGTCTTGGGCATGTGCACCATG
TTCATCGTAGTCTGTGCCTTGGTTGGAGTATTAGTGCCAGAGCTCCCTGACTTCTCTGATCCATTGCTGGGTTTT
GAACCAAGAGGAACAGCAATAGGCCAGAGATTGGTCACATGGAATAATATGGTGAAAAATACAGGATACAAAGCA
ACATTAGCAAATTATCCCTTTAAATATGCAGATGAACAAGCCAAAAGCCATCGGGATGATAGATGGTCAGATGAT
CATTATGAAAGAGAGAAAAGAGAAGTTGACTGGAACCTCCACAAGGACAGCTTTTTCTGCGACGTTCCAAGTGAC
CGATATTCCAGAGTGGTATTTACTTCATCTGGAGGGGAGACATTATGGAATTTACCTGCAATTAAATCAATGTGC
AATGTAGATAATTCCAGGATCAGATCTCATCCCAGTTTGGTGATCTCTGCCAGAGGACCACTGCTGCCCTCCTGC
TGCCCCAGCTGGACACTGGGAAACTACATCGCCATTCTGAACAATAGATCGTCCTGTGAGAAAATAGTTGAGCGA
GACGTTTCTCATACCTTGAAGCTGCTTCGGACTTGTGCCAAACACTACCAAAATGGCACTCTGGGGCCAGACTGC
TGGGACATGGCAGCCAGAAGAAAGGACCAGCTCAAGTGCCACCAATGTGCCACGCAAATGTACCAAGTACAATGCT
GTGTACCAGATCCTCCATTACTTGGTGGACAAAGACTTTATGACCCCAAAGACGGCTGACTATGCCACGCCAGCT
TTAAAATACAGCATGCTCTTCTCTCCACAGAGAAAGGGGAGAGCATGATGAACATTTACTTGGACAACCTTTGAA
AACTGGAACCTTCTGACGGCGTGACTACCATCACCGGATTGAGTTTGGTATCAAACACAGTTTGTTCAGGAT
TATCTTCTAATGGATACTGTGTATCCTGCCATAGCCATTGTGATTGTCCTTTTAGTTATGTGTGTCTACACCAAG
TCCATGTTTATCACTCTGATGACAATGTTTGCAATAATCAGTTCTTTGATTGTTTCTTAAAAAAAAAAAAAAAAAAAA
A

653/6881

FIGURE 607

MAMSNNGNDFVVLNSSSIATSAANPSPLTPCDGDHAAQQLTPKEATRTKVSPNGCLQLNGTVKSSFLPLDNQRMF
QMLPQCCHPCPYHHPLTSHSSHQECHPEAGPAAPSALASCCMQPHSEYSASLCPNHSPVYQTTCCCLQPSPSFCLH
HPWPDHFQHPVQQHIANIRPSRPFKLPKSYAALIADWPVVVLGMCTMFIVVCALVGVLVPELPDFSDPLLGFEP
RGTAIGQRLVTWNNMVKNTGYKATLANYPFKYADEQAKSHRDDRWSDDHYEREKREVDWNFHKDSFFCDVPSDRY
SRVFTSSGGETLWNLPAIKSMCNVDNSRIRSHPOFGDLCQRTTAASCCPSWTLGNYIAILNNRSCQKIVERDV
SHTLKLRTCAKHYQNGTLGPDWCWMAARRKDQLKCTNVPRCKTKYNAVYQILHYLVDKDFMTPKTADYATPALK
YSMLFSPTEKGESMMNIYLDNFENWNSSDGVTTITGIEFGIKHSLFQDYLLMDTVYPAIAIVIVLLVMCVYTKSM
FITLMTMFAISSLIVS

654/6881
FIGURE 608

GCTTCCGCCTCGGCTCCTCTTCTGCGGGCATCCGGGATCCCTACGTCCCGCGTCCCCGAGCGCTCGGAGCCTA
CGCGCCCAGCGCTACCGAAACCCAGAGTCTTGCGCCCTGGAGTCCCCGCGCCCCGAGCCCCGAGCACCCGGGAGT
CCCCGAGCCTCGCGCCCCGGAGTGCCCGAGCCTGCGCCGCGCGCACCCCGGATACCCCGGGTCCCCGCGAGCTGCCGA
GGCCGCCCCGCGCGCCCCGCGGGACAGTACCGCCTTCTCCCTCTGTCCGCGCCATGGCCGCCCCCGACCTGTC
CACCAACCTCCAGGAGGAGGCCACCTGCGCCATCTGCCTCGACTACTTCACGGATCCGGTGATGACCGACTGCGG
CCACAACCTTCTGCGCGAGTGTCATCCGGCGCTGTGCGGGCCAGCCCCGAGGCCCTACGCGTGCCCCGAGTGCCGC
GAGCTGTCCCCGAGAGGAACCTGCGGCCAACCGCCCCGCTTGCTAAGATGGCCGAGATGGCGCGGCGCCTGCAC
CCGCCGTGCGCGGTCCCCGAGGCGTGTGCCGCGCACCGCGAGCCACTGGCCGCCCTTCTGTGGCGACGAGCTGCG
CCTCCTGTGTGCGGCCCTGCGAGCGCTCTGGGGAGCACTGGGCGCACCGCGTTGGCCGCTGCAGGACGCGGCCGAA
GACCTCAAGGCGAAGCTGGAGAAGTCACTGGAGCATCTCCGGAAGCAGATGCAGGATGCGTTGCTGTTCCAAGCC
CAGGCGGATGAGACCTGCGTCTTGTGGCAGAAGATGGTGGAGAGCAGCGGCAGAACGTGCTGCGTGAGTTGAGC
GTCTTCGCGCTTTGCTGGCAGAGGGAGGGACAGCAGCTGCTGCAGAGGCTGGAGAGGAGGAGCTGAAGGTGCTGC
CCCCGTGCGGGAGGCGCAGCCCACCTAGGCCAGCAGAGCGCCCACCTAGCTGAGCTCATCGCCGAGCTCGAGAG
GCCGTGCCAGCTGCCTGCGCTGGGGCTGCTGCAGGACATCAAGGACGCCCTGCGCAGGGTCCAGGATGTGAAGC
TGCAGCCCCCAGAAGTTGTGCTATGGAGCTGAGGACCGTGTGCAGGGTCCCCGGGACTGGTAGAGACACTGCGGA
GGTTTCGAGGGGACGTGACCTTGGACCCGGACACCGCCAACCTGAGCTGATCCTGTCTGAAGACAGGCGGAGCG
TGCAGCGGGGGACCTACGGCAGGCCCTGCCGGACAGCCCAGAGCGCTTTGACCCCGGCCCTGCGTGCTGGGCC
AGGAGCGCTTACCTCAGGCCGCCACTACTGGGAGGTGGAGGTGGGGACCGCACCACTGAGCTGGGCCCTGGGGGTGT
GCAGGGAGAAGCTGAACAGGAAGGAGAAGGGCGAGCTGTCCGCGGGCAACGGCTTCTGGATCCTGGTCTTCTTG
GGAGCTATTACAATTCTCGGAACGGGCCCTTGGCTCCACTCCGGGACCCACCCAGGCGCGTGGGGATCTTTCTGG
ACTACGAGGCTGGACATCTCTTTCTACAGTGCCACCGATGGGTCACTGCTATTATCTTTCCCGAGATCCCT
TCTCGGGGACGCTGCGGCCCTCTTCTACCCCTGTCCAGCAGCCCGACCCGATGACTATCTGCCGSCCGAAAG
GTGGGTCCGGGGACACCTGGCTCCCCAGTGACTCGGGCCCTCCTGGAGGAGTCTGTGCTCTCTCTGCCCTC
CAGGCCACTGAGTGTGTTTGGCCACTTGGAGGACCTGGGAGGAGGGAGTGTGCTTTGAGCAAGAGGAGGAATC
CTGGTGCTTTCTGAGCCTGCGTGGGAGAACCCCAATTCTAGCACTCCAGGAACTGTGGGAGAGTGTGGGGCAG
GCTCCGTCTCTCCCTGGGAGACCCCTCCAGCCACCGGGTGCCACTTAATGCCAACAGCCCTTACCAAAGCTGGGAG
CCCCATTGCCCCGGCAGCTCTGGCCTGTGGTTCCAGAAGCTGAGAAAATCCACTGGGGCTTGAGAAATCCAGGG
TTCACCTAAGCTGCACAGTTCTGACGCTTTGTGAGCCCCCTGAAAGTCTTGTGTACCCACCTCTGAAGATGCT
GGGGAAGGCAGCTGGGATGGGAGCCAGCCCATGCCTGTCTGTGACCCACAGTGGGTGAGAGCCCGTCACAGTC
TGAGGACAGGAACAGAGAGTAGATGTCTATAATAAAGGGGCTTCTGGGAGGTGCTGGGCACAGATGTCTGTTC
AGCAGGTGTGTGGGCTAGAGGAGAGAGCAGAGCCAGAAATGTCTTTTGAGGCCACGTTCTGACTTGAAGCT
TTCGTGGGCATGTTGCCATTGGGTTTTGCCCTTGCAAAGGCTTCTAGGTCTCAGTGGGCCCTCAGGACCCAGG
GTCCAGCTGCTGCTTGGGGATGTGCACTGCTGCGCCGCGCTTGCAGTCTCTTACCCTGGGGAGGAACAGTGGC
TTCTCAGAGCCTGGGGCATACAGAAGAAGGCAGGAGTTGATTTTGTGTTGGGTTTGGGGTTTCTTTGTCTCAA
GGTACTGTTCTGTTTCTTTACCCCTCTGCTTATTTATTGTAAGCATTCCACGTTAAATAAACTTTGGCTGT
TGTCTAC

655/6881
FIGURE 609

MELRTVCRVPGLVETLRRFRGDVTLDPDTANPELILSEDRRSVQRGDLRQALPDSPERFDPGFCVLGQERFTSGR
HYWEVEVGDAAGPWGCAGRT

656/6881
FIGURE 610

CCGCAGGCAGTAGAGTGAAGCGCGCAGCTGCCAGGACTTGCGCGGTGACGTGCGCCGCTGCCAGGACCTTGCAGG
TGGAGAGCATAGTTGCCAAAATCAAGGCGGAGGAGCGCACCGCCGCTAGGATCCAGGCGGAGAAGCCCACCGCGG
CCAGGACCTAAGGATGCAGTACACTGCTGCCAGGATCTTGTCTGTGGAGCGCAGCGCGGCCAGGACCTCCGGCTG
CAGCACACCGCTGCCAGGATCTTATCGGCAGAGCGCTCCGCGGTCCGGACCCCGCCCCGTGCGCGTCCCCGACCC
CGCCCCGTGCGCGTCCCCGGCGTTGGCGTCTTCGTCTCTGTTGCTGGTCTCCGTCCGGTCCGGCGCGTCTAGGTC
TCCGGCCCTCCCCAGCCGCTCCTGCGCCCTTGCCGGCCCCGCGCCCGCAGCCCTGGCGCTCCCTGCGGGCCCCG
CCGAGGCCCGCTGCGCCCTGTGCCAGCGCGCGCCCCGGGAACCGGTGCGCGCCGACTGCGGCCACCGCTTCTGTC
GGCGTGTGCTGGTGCCTTCTGGGCCGAGGAGGACGGGCCCTTCCCGTCCCCGAGTGCGCCGACGACTGCTGGC
AGCGCGCCGTGGAGCCCCGCGCAGGCCCGCTCAGCCGCGCCTTCTGGCGCTCGAGGAGGCGGCGCGCGCCCCG
CGCGCAGCGGCCCGGCCAGCGAGGCCGCGCTGCAGCTGCTGTGCCGCGCCGACGCCGGCCCGCTCTGCGCCGCT
GCCGTATGGCTGCGGGCCCCGAGCCGCCGAGTGGGAACCGCGCTGGAGGAAGGCGCTGCGCGGAAGGAGAACA
AGGGGTCTGTGGAATCATGAGAAAGGACTTGAATGACGCCCGGGACCTGCATGGCCAGGCAGAGTCAGCAGCTG
CAGTGTGGAAGGGACACGTGATGGACCGTAGGAAGAAGGCACTGACCGACTACAAGAAGCTGCGGGCCTTCTTTG
TGGAGGAGGAGGAGCATTTCCTGCAGGAGGCTGAGAAGGAGGAGGGGCTCCCTGAGGACGAGCTGGCTGACCCCA
CTGAGCGGTTTCAGGTCACTGCTGCAGGCGGTCTCGGAGCTGGAGAAGAAGCATCGCAACCTGGGCCTCAGCATGC
TGCTGCAGTGATGGCGCAACCCGTGGCAGTCCCAGAGCTGGAGGCAGGAGGATGGATCCTCATCTCCATGGGAA
GTGTCAGCGTGTGGCTGCCAGGGAAGCGTGGCAGGCGCCTGGCCTTGGGTCCATCTACATAGTTGCGTGTTCAA
CAATGTCCATTTATCCTTCACCCGAGGCGTGTTCCTGGGGGCTGCAAACACCTCCCTGTGCTCCACCTGCCTCCG
CAGAAGGAAGCCTCTTTCTCTGTTTCCTGGGTGAGGGGGCTGGCAGGTGGCTAACCCATTTAGCATCTCCAGG
CCCTGCCATCGTGTCTCATCTTGCTGTTATCTCTAGCTCTTCCCTCCTCCATTTCTTTAGTAGTTGAATTTT
GCAAAGCTTGTAGCAGTAGCTCAGTTGCCTGCAGCATCCTTGTGTGTAGATAAATTAGTCGACAGAACTCAGCA
CTGGGGACAGGATTGCAAAGTCGGGGACATAGATGCAGACAGTTGTTGAGATTGGGGATAGCCGGGCTTGTGAG
CGGTGCCCCATTTCCAGATGAAGCCTTTCAGCCCTTCTGAGTCCCCGGCCCTTGGTGCATGTCTGTGAGTTTGAC
CTGCCCAGCGTGTGGGCTGGCTCAATGCTGAATAAAGTGGGTTTGTGTGAGCTCGTTTGCTTCGTCTCCGTGTGT
CCACCTGGCCTCTTCCCCCTGCCCTGGCCACCTCCAGTGTCAAAGGAACTTCCTCGTGACACGTGCTAAAGCA
TGGTGAGGAGGACTTTGATTGGGACCATGAGATGGGTGTGGGACCCTTTCCTTGGGGCCTGGGGGGAGATGGGG
CTCCACCCGACGTAGCAGGGCAGGGTTGGAGGAGCAGGAGCAGTATAGGGTCCATGGGTGGGAATGACTGTG
AGGAGACATCAGGGCTGAGGGGGCTCTGGCTAAACCCACCTCACAGAGTCCTTGCTGCAGGCAGGCAGGGCGATC
AGA

657/6881
FIGURE 611

MQYTAARILSVERSAARTSGCSTPLPGSYRQSAPRSGPRPVRVPDPAPCASPALASSCCWSPSGRRPSRSPALP
SRSCALAGFAARSPGAPCGPRRGRLRPVPARAPGTGARRLRPPLLSGVRGALLGRGGRALPVPRVRRLLAARRG
ARQAPAQPPPSGARGGGRGARARRPGQRGAAAAVPRRRRPA LRRLPYGCGPRAARVGTALEEGAARQGEQGVCG
NHEKGLE

658/6881
FIGURE 612

GGCACGAGGCGCTGGCCGGGCTGGCTGCGGCCGCCCTGGGCCGGGCCCCACCGGACGGCTTGCTCTGCTCTTTAC
CTGGGGTTGCTGTGCGAGGACCCTGTGCAAGACTCGGCCGGTTTTTCTTTCTCCCTGATGGACAGACCCAAACATA
GCCGCGCAGCATCGTGAAGGGCTGGGGCCTTCACTCCTCTGTGGCTCTGGAAGAGCCCGATTTCCTCAGGAGGCA
TGTCGGGCCCCAGGCCTGTGGTGCTGAGCGGGCCTTCGGGAGCTGGGAAGAGCACCCCTGCTGAAGAGGCTGCTCC
AGGAGCACAGCGGCATCTTTGGCTTCAGCGTGTCCCATACACGAGGAACCCGAGGCCCGGCGAGGAGAACGGCA
AAGATTACTACTTTGTAAACAGGGAGGTGATGCAGCGTGACATAGCAGCCGGCGACTTCATCGAGCATGCCGAGT
TCTCGGGGAACCTGTATGGCACGAGCAAGGTGGCGGTGCAGGCCGTGCAGGCCATGAACCGCATCTGTGTGCTGG
ACGTGGACCTGCAGGGTGTGCGGAACATCAAGGCCACCGATCTGCGGCCCATCTACATCTCTGTGCAGCCGCCTT
CACTGCACGTGCTGGAGCAGCGGCTGCGGCAGCGCAACACTGAAACCGAGGAGAGCCTGGTGAAGCGGCTGGCTG
CTGCCCAGGCCGACATGGAGAGCAGCAAGGAGCCCGGCCTGTTTGTATGTGGTCATCATTAAACGACAGCCTGGACC
AGGCCTACGCAGAGCTGAAGGAGGCGCTCTCTGAGGAAATCAAGAAAGCTCAAAGGACCGGCGCCTTGAGGCTTGC
TGTCTGTTCTCGGCACCCCGGGCCCATACAGGACCAGGGCAGCAGCATTGAGCCACCCCTTGGCAGGCGATACG
GCAGCTCTGTGCCCTTGGCCAGCATGTGGAGTGGAGGAGATGCTGCCCCTGTGGTTGGAACATCCTGGGGTGACC
CCCGACCCAGCCTCGCTGGGCTGTCCCTGTCCCTATCTCTCACTCTGGACCCAGGGCTGACATCCTAATAAAAT
AACTGTTGGATTAGAAAAAAAAAAAAAAAAAAAA

659/6881

FIGURE 613

MSGPRPVVLSGPGAGKSTLLKRLQLQEHSGIFGFSVSHTRNPRPGEENGKDYYFVTREVMQRDIAAGDFIEHAE
FSGNLYGTSKVAVQAVQAMNRCVLDVDLQGVNRNIKATDLRPIYISVQPPSLHVLEQRLRQRNTETEEESLVKRLA
AAQADMESSEKPGFLFDVVIINDSLDQAYAEELKEALSEEIKKAQRTGA

660/6881
FIGURE 614

GGCACGAGGTCGGGCCTGGGTCCGACGGTAGTGGGTAGCGGGTCTCGGGTTGCGGGTTGCAGGTTGCAAGCCGCA
GGCCCCAGGCAACTGCCCTCCCGGCGCCATGTTTCGGCTCCAGTCGTGGAGGCGTGCGCGGCGGGCAGGACCAGTT
CAACTGGGAGGACGTGAAGACTGACAAGCAGCGGGAGAATACTGGGCAACTCGCTGATGGCGCCGGTAGGCCG
CTGGCAGAAGGGCCGCGACCTCACCTGGTACGCCAAGGGCCGGGCGCCATGCGCGGGCCCGAGCCGCGAGGAGGA
ACTGGCAGCCGTGCGGGAGGCGGAGCGCGAGGCGCTGCTGGCCGCCCTTGGCTACAAGAACGTGAAGAAGCAGCC
CACGGGCCTGAGCAAGGAGGACTTCGCGGAGGTCTGCAAGCGGGAAGGAGGCGACCCCGAGGAGAAGGGCGTGGA
CCGGCTGCTGGGGCTGGGGAGCGCAAGTGGCTCCGTGGGCGCGTGCGCATGTCCCGAGAGGACAAGGAGGCCGC
CAAACCTGGGGCTGTCTGTGTTACGCATCACCGCGTAGAGAGCGGCGGGCCCGGGACCTCGGCAGCCTCGGCCAG
GAGGAAGCCCGGGCGGAGGATCAGACGGAAGCAGTTGTGAGAGCCACAGGAAAAGCAAGAAGGAGAAGAAGAA
AAAGAAAAGAGGAAACACAAGAAAGAGAAGAAGAAGAAAGACAAAGAGCACAGGCGGCCAGCTGAGGCCACCTC
CTCTCCACATCTCCTGAGAGGCCAGGCACCACCACCATGACTCCGACTCCAACCTCCCCCTGCTGTAAGAGGAG
GAAGCGGGGACACAGTGGGGACAGGAGGAGCCCGTCTCGCAGGTGGCATGACAGAGGCTCTGAGGCCCTGATGGCT
GGACCCTGCTCACTGCTGTTGTGGGACCCTGAACCCTCCCTTACCTTGCTTGCTCCTGCTCGGAAGCTCCTT
GGGTGTGGGTGAAGCCCGAGGCTGCTCCTGTGGAAGTGGCTCTGGGCACCAGCCTGTGGGGCTAAAGACTTGACA
GCTAGCTCTGGAGCAGCCGGCTTCCTGGAAAACCTCCAGGTTTCGCATACCAGGGATGGCCCTGGCTTGGCCTG
CGAAGGTGAACCTGCCCAGATTTATCAGTAGAGGCTGGACTCCCTCIGTGTCTGCTGCCCATGGTTGCAGCAGCCAT
GGGCCTATGAGCGGTCTAACTGTGGCCAAGTATGGTGACCTCTATTTTCTTTATATTGACTCTTTGTATTTCAA
TAAATATATTTTAAAGGAAGGTAAAAA

661/6881

FIGURE 615

MFGSSRGGVRRGGQDQFNWEDVKTDKQRENYLGNSLMAVGRWQKGRDLTWYAKGRAPCAGPSREEELAAVREAER
EALLAALGYKNVKKQPTGLSKEDFAEVCKREGGDPEEKGVDRLLGLGSASGSVGRVAMSREDKEAAKLGLSVFTH
HRVESGGPGTSAASARRKPRAEDQTESSCESHRKSKKEKKKKKKRKHKKKKKKDKEHRRPAEATSSPTSPERPR
HHHHDSDSNSPCCRRKRGRHSGDRRSPSRRWHDRGSEA

662/6881
FIGURE 616

AGGAAGCCCTGCTTGATGAGCATGAGGCTCCACGGGGCAGTGGCTGTGACGGTGAGTGCCACACAGAGCTGCCC
ACACTGGTGGAGGGAAGGCAGGGAGATACCAGGACCATCCGGAAGGGGCTGAGTGTCAATTTGACAGGTGCCATGT
GAGCAGAGATGTGAAGGAGTGGCCCGGGACAATCAGGGCAGAATCCCTGAGGTGTCCCTGGCCAGTGTCTTCCA
CCTGTCCACAAGCATGGGGAACATCTTCGCCAACCTCTTCAAGGGCCTTTTGGCAAAAAGAAATGCCGATCCT
CATGGTGGGCTGGATGCTGCAGGGAAGACCACGATCCTCTACAAGCTTAAGCTGGGTGAGATCGTGACCACCAT
TCCCACCATAGGCTTCAACGTGGAAACCGTGGAGTACAAGAATCAGCTTCACTGTGTGGGACGTGGGTGGCCA
GGACAAGATCCGCCCCCTGTGGCGCCACTACTTCCAGAACACACAAGGCCTGATCTTCGTGGTGGACAGCAATGA
CAGAGAGCGTGTGAACGAGGCCCCGTGAGGAGCTCATGAGGATGCTGGCCGAGGACGAGCTCCGGGATGCTGTCTT
CCTGGTGTTCGCCAACAAGCAGGACCTCCCCAACGCCATGAATGCGGCCGAGATCACAGACAAGCTGGGGCTGCA
CTCACTACGCCACAGGAACCTGGTACATTCAGGCCACCTGCGCCACCAGCGGCGACGGGCTCTATGAAGGACTGGA
CTGGCTGTCCAATCAGCTCCGGAACCAGAAGTGAACGCGACCCCCCTCCCTCTCACTCCTCTTGCCCTCTGCTTT
ACTCTCATGTGGCAAACGTGCGGCTCGTGGTGTGAGTGCCAGAAGCTGCCTCCGTGGTTTGGTCACCGTGTGCAI
CGCACCGTGCTGTAAATGTGGCAGACGCGAGCCTGCGGCCAGGCTTTTTATTTAATGTAAATAGTTTTTGTTC
ATGAGGCAGTTTTCTGGTACTCCTATGCAATATTACTCAGCTTTTTTTATTGTAAAAAGAAAAATCAACTCACTGT
TCAGTGCTGAGAGGGGATGTAGGCCCATGGGCACCTGGCCTCCAGGAGTCGCTGTGTGGGAGAGCCGCCACGC
CCTTGGCTTTAGAGCTGTGTTGAAATCCATTTTGGTGGTTGGTTTTTAACCCAACTCAGTGCATTTTTTAAAT
AGTTAAGAATCCAAGTCGAGAACACTTGAACACACAGAAGGGAGACCCCGCTAGCATAGATTGCAATTACGGC
CTGGATGCCAGTCGCCAGCCCAGCTGTTCCTCGGGAACATGAGGTGGTGGTGGCGCAGCAGACTGCGATCAAT
TCTGCATGGTCACAGTAGAGATCCCCGCAACTCGCTTGCTTGGGTACCCCTGCATTCCATAGCCATGTGCTTG
TCCCTGTGCTCCCACGGTTCCCAGGGGCCAGGCTGGGAGCCACAGCCACCCCACTATGCCGAGGCCGCCCTAC
CCACCTTCAGGCAGCCTATGGGACGCAGGGCCCCATCTGTCCCTCGGTGCGCGTGTGGCCAGAGTGGGTCCGTG
TCCCCAACACTCGTGCTCGCTCAGACACTTTGGCAGGATGTCTGGGGCCTCACCAGCAGGAGCGCGTGCAAGCCG
GGCAGGCGGTCCACCTAGACCCACAGCCCCCTCGGGAGCACCCACCTCTGTGTGTGATGTAGCTTTCTCTCCCTC
AGCCTGCAAGGGTCCGATTTGCCATCGAAAAAGACAACCTCTACTTTTTCTTTGTATTTTGATAAACACTGAA
GCTGGAGCTGTAAATTTATCTTGGGGAACCTCAGAACTGGTCTATTTGGTGTGCTGGAACCTCTTACTGCTTT
CAATACACGATTAGTAATCAACTGTTTTGTATACTTGTTCAGTTTTTCATTTGACAAACAAGCACTGTAAATTA
TAGCTATTAGAATAAAATCTCTTAATATT

663/6881
FIGURE 617

MGNIFANLFKGLFGKKEMRILMVGLDAAGKTTILYKLKLGEIVTTIPTIGFNVETVEYKNISFTVWDVGGQDKIR
PLWRHYFQNTQGLIFVVDSDNRERVNEAREELMRMLAEDELRDVLLVFANKQDLPNAMNAEITDKLGLHSLRH
RNWYIQATCATSGDGLYEGLDWLSNQLRNQK

664/6881
FIGURE 618A

GTGGGACCATCCAGTTGCAGGAAAACAAGCTTAACACGCCCACTGATTCTACATTATGCTCCTACCTCCCAGCAG
CCTCTCCAGGCCCAGAACTTTCTCCAGTCAGCCTCTACAGACCAAGCTCATGACTCACAATGGCCTATTTAGGCC
CATACCTTACCTCAGGCAGTCTCCGCAGATGAGCCTACTGCCTCACAACAGCCTCCACAGGCACAGCTCCATCG
TTACAATGGCCTCTTTAGACCCAGCTCCTGCCTCCCAGCCTTCTCTCCAGGCCCTGAACTTTCTCAAGTCGACCT
CACCAGGCCCAGCTCATGCTTCTTTGCAGCCTCTCCAGGCCCAGCTCCTGCATCTTGGTGGCACCTCCAGGCCCA
GCCTCTGCCTCCCGTCGGCCTCTACAATCCCAACATCTGCCTCACAGCAGATTCTTCAGGCCCAGCATCTGCCTC
ACTGTGGACCCCCAAGCCAAGCTCCCAACCTTTTCAGCAGCTTCTACACACCCAACCTCCTGCCACCCAGTGGCCT
CTTTAGGCCAAGCTCATGCTTACAAAGGGCCTTTCCAGGCTCAACTTTTGTCTCATGGCAACCTTCTCTGGCCAG
ATTCTGCCTGTCTCCACAGCAGCCTAGACAGGCCCAGGCTTGCCTCACACTGGCCTCTCTACATCCAGCTCATG
CCTCACGGTGGCCTCTCCAGGCCCAACTCCTGTCCAGGACGTCTCTCCGGGCCCAAACTTACTCAAGTCAGA
CTCTCTAGTCCCAACTGTCTGCCTCCTGGTGGCCTATGAAGGCCCAAAATCTCCTCAAGTGGACCTCTCCAGGCC
AGCTCCTGCCTCCTGTCTCAGCGTCTACAGGCCCAACCTCTGCCTCATGGGGCTTCTCCAGGCCACCTCTTCTC
TTGGCTGGGTCTACAGGCACAAGTGTCTGCCTCACAACAGCCTTTTTTGGGCCAGTTCCTGTCCAGCTCATGGCGG
CCAATGTAGGCCCAAACTTCTCCTCAAGTCAAACTCTCCAGGCCACCTTCTGCTTCCCGGTGGCATGAACAGGCC
CAGCTTTGACTTGAGAACAGCCTCTGCAGGCCCTGTCTTGCCTCCCAGGGGCTTTTCCAGACCCAGCTCCTGC
CTCATGGCAGCTGCCCCAGGCCAAATTTCTGCCTGCCTGCCAGCAGCCTCAACAGGCACAGCTCCTCCCTCACAG
TGGCCCATTTAGGCCCAACTCATGACTGTCCGGCCATTTCCAGGCCTAGTGCCTGCCTCGTGGCTGACTCTTGAA
GCCCAAACTTCTCCTCAATCAGCCTTTTGGCCAACTTCTGTCTACTGTCTCGACTCTACAGGCCAGCCTCTGCCTC
ACAGTGGACCTTCCAGACCCAGATGGTGTCTCACTGTGGCATCCTCAGGCGAAGCTCCTGCCTTTCCGCAGCCTC
TACAGGCCCAGCTCCTGCCTTGCAGTGGCCTCTTTAGGCCAAGCTCATGCCCCATGGCGACTTTTCCAGGCCAG
CTTTTGCCTTTTGCAGCCTGTGCAGGCCCAGAATGTCTTAACTCGGCATCTCCAGGATGAGCTCATCCTCCAG
TGCGTCTACAGGCCCGTCTCCTGCCTCACAACAACCTCCTTTGGCCCAACTCCTGCTGAGCTGTGGCAGCCTCT
GTAGGCCACAGAATTCTTAAGGTAAAGCTTTCCAGGCCACCTTTGGCCTCCCGGCAGGCTCAGCAATCAAACTA
TTCCCTCACTGCGGCCACCGAAAGCCAAGTTTCTCCCTGCCTCACGGCATCCTCCGAAAACCTGAGCATTTGCCTC
ACGGTGGCCTCCCCAGGCCATGAATCTGCCTGCCTCCCAGGCAGCTGTGCCTCACAATGGTCTCTTTAGGCCCA
GCTCATGTCTAAAGATGGACTCTCCAGGCACAGCTCTTGCCTCCTGGCAGCCTCTGCAGGCCCAATTTCTCCAA
AGTTGGCCTCTCCTAACTCAGCTCCTGCCTCATGTCTGCCTACACAGGCCCAGACTCTTACCACACAGTAGACCC
TCCAGGCCCACTTGCCTGATCATAGCCTCTTAAAGGCCAAGCTCCTGCCTTTCCGGCAGCCTCTACAGGCCAAG
CTCCTGCCTCGCAATTGCCTCTGTAGGCCAAGATCATGCCGTGAAGTGGCCTTCCCTAGCCTAACTTTTGCCTTT
TGACGCATACTCCAGTCCCAAACTTCTCCTCAGTCAGCCGGTCCAGGCCAAGCTCTTCTCCCAAGGCTTCTGC
AGGCCAAAATCATCCTGAAGTCAACCTCTGCAGGCCGAGCTCCTGCCTCCAAGTGTGTGTAGGCCAAGCTAATG
CCTCACAGCACACTTTCCAGGCTGAGCATTTCTTTTGTGCATCCTCTCCAAGCCCTGAACCTTACTCCAGTTGGC
CTCTCCAGACCAAGCTCTCCCTCCAGTGGCCTCTACAGGCCAAAATGTCTCAGGTGAGCCTCTCCAGGGCCA
ACTCCTAGCTACCGGTGGCTTCTGCAGGCCAAAATCGACCTCAAGTCAGCCTCTTACACCCAGCTCTTGCCTCT
GAGTGGCCTCTCCAGGAGCAAACTTCTCAAGTCGGCCTCTCCAGGCCCAGCCTCCTGCTTCCCGAGGGCATGT
ACAGGCCCAGCCTCTGCCTCACAGCAGACTCTTACACCCAGCTCTTCCCTGTCTGCGGCCTCTCCAGTCCAAAG
CTGCTCCTGCCTTTTGGCAGCTTGTACAGGCCCAGCTCCTCCCTCACGGTGGCCTCTTTCCGGCCCAACTCATGCC
TCTTGCAACGTGCCCAAGTGTAGCTCCTGCCTCACACTGGCCTGTTGAGGCCCAGCTCATGCCTCTCGTGGCCT
CAACGGGCCCATCCCCTGCCTGTGCGCGGCCTCTACAGGCCCGGCCTCTACCTCACAGTGGGCTCTCCAGGCCCA
CCTCTTCTCTACCGTGGCCTCCTGGGCAATGCTCTCTCCCTCTCGGGAGCCTCTGTGGGCCCAGCTCCTGCCTCC
CAGTGGCCTCTGCTGGCCAAAGCCCGTGCCTCAGGGCAGCCTTTCCAGGCCTAGCGTTTGTGCTTTGCATCCTCT
CCAGGCCCTGGACTTCTCCAGTCCGGCTCTCCAGGCCCAGCTCTTCTCTCGGCGGCCTCTGCAGGCCCAGACT
GTCGTCAAGTCGGCTGTCTCCAGGGCCAGCTCCTGCCTCCCGGCGGCCTCTGCAGGCCCAAGTCGTCTCAAGTTG
GCCTCCCCAGGCCCAGCAACGGCCTCTCGGCGGCCTTTCCGGGTGCAAAAGTTCTCGAGTCAGCCTCTCCAGGC
CCAGCTCCTCCTGCCTCCCAGTGGCCTCTTTCCGGCCAGCCAGCTCATGCCTCCCGGCGGCCTTCCAGGCCCT
GCTTTTGACTTTTGGTGGCCTCTGCAGCCTCGACAAGGCCAGGCCTCCTGCCTCCCAAGGCCTGCACAGGCC
AGCCTCTGCCTCACAGCAGACTCTCCACGCCCAGCTAGCTCTGCCTCACTGCGGCTTCCCGAGTCCAAAGCTCC
TGCTCTCAGCCGCTTCCGCAGGCCCAGCTCCCGCCTGCCAGTGGCCTCTTCCAGGCCATGGGGCTCATTTCTCTCA

665/6881
FIGURE 618B

CAACGGCCTTTCCAGGCCAGTTTTCCCTTCCGGCGGCCTCTTGGCCTCTAATTGTTTATCTTTGGGTATAA
ATCCCAAATATTGAATTTGGAATATTCCACCATT

666/6881
FIGURE 619

TCTAATACCTATTGATCTGTCACTTTCTCCCATCACGCTCAGGTGGGACCATCCAGTTGCAGGAAAACAAGCTTA
ACACGCCCCTGATTCTACATTATGGTGAGTTCTATAATTATTTTATTATATATTACAGTGTAATAATGGAAATA
AAGTGCCTAATAAATGTAAATGTGCTTAAATCTTTTGGCCCAGCTCCTACCTCCCAGCAGCCTCTCCAGGCCAG
AACTTTCTCCAGTCAGCCTCTACAGACCAAGCTCATGACTCACAATGGCCTATTTAGGCCCATACCCTACCTCAC
GGCAGTCTCCGCAGATGAGCCTACTGCCTCACAACAGCCTCCACAGGCACAGCTCCATCGTTACAATGGCCTCTT
TAGACCCAGCTCCTGCCTCCCAGCCTTCTCTCCAGGCCCTGAACCTTCTCAAGTCGACCTCACCAGGCCAGCTC
ATGCTTCTTTGAGCCTCTCCAGGCCAGCTCCTGCATCTTGGTGGCACCTCCAGGCCAGCCTCTGCCTCCCGT
CGCCTCTACAATCCCAACATCTGCCTCACAGCAGATTCTTCAGGCCAGCATCTGCCTCACTGTGGACCCCCA
GACCCAGATGGTGTCTCACTGTGGCATCCTCAGGCCAAGCTCCTGCCTTTCCGCAGCCTCTCCAGGCCAGCTCC
TCCTGCCTCCCAGTGGCCTCTTTCCGCCAGCCCAGCTCATGGCTCTCGGCGGCCTTCCCAGGCCCGCTTTTGA
CTTTTGGCGGCCTCTTCAGGCC

667/6881

FIGURE 620

TTGAAGAACTTTGCCAAATACTTTCTTCACCAATCTCATGAGGAGAGGGAACATGCTGAGAAACTGATGAAGCTG
CAGAACCAACGAGGTGGCCGAATCTTCCTTCAGGATATCAAGAAACCAGACTGTGATGACTGGGAGAGCGGGCTG
AATGCGATGGAGTGTGCATTACATTTGGAAAAAAGTGTGAATCAGTCACTACTGGAAGTGCACAAACTGGCCACT
GACAAAAATGACCCCCATTTGTGTGACTTCATTGAGACACATTACCTGAATAAGCAGGTGAAAGCCATCAAAGAA
TTGGGTGAGCACGTGACCAACTTGTGCAAGATGGGAGCGCCCGAATCTGGCTCGGCGGAATACCTCTTAGACAAG
CACACCCTGGGGGACAGTGATAATGAAAGCTAAGCCTCAGGCTAATTTCCACATAGCCGTGGGAGTGACTTCCCT
GGTCACCAAG

668/6881
FIGURE 621

MKLQNQRGGRIFLQDIKKPCDDWESGLNAMECALHLEKSVNQSLLELHKLATDKNDPHLCDFIETHYLNKQVKA
IKELGEHVTNLCKMGAPESGSAEYLLDKHTLGDSDNES

669/6881
FIGURE 622

ATGGCCGGCACAAAGATGGGGCGGGAGGACAGCGTGCTGGATACGGCGTGGGGCGGGTCCGCGCCCCGGGAGACC
CATGCTGGGGCGGCCCTGTCACCTCGTCTCGCTGCCAAGCACAGGGCTGAGGTGCGGAGGGACCACTGAGAGGACT
TCGCTGCGTGTCAGCTCAGCTCTGGACGGCGTGGAGGAGCCCTGGAACTTTATTTACAAAACTGACAGCAGCT
GCCCTTAGTTTGGCAGCTGGAAAATGGGAAAGAGAAAGTAAGAGGAAGGGTGGCAAAGGAGGGAGCCCATGTGGA
CAGCAGGAACGGGTACACGATAGCACAAGAATCAAAACAAAGGCTCTCTGGTCGGAGTTTAGAAAGGCTGAAGGG
GATACTAAAGGAGGAGATAAAGCCCAGGTGAAGGACGAACCACAGAATAGATCCGCGAGGTTGTCTGCTAAACCT
GCTCCTCCAAAACCAGAGCCCCAAGCCTAAAAAGGCCCTGCAGAGACAGTACCCAAAGGGAAAAAGGGAAAAGCT
GTTGCTGGCAAGGAGGGGAATAACCTGTCAGAAAATGGAGATGCCAAAACAGACCAGGCACAGAAAGCTGAAGGT
GCTGGAGATGCCAAGGCAAATCTGCTGCCACGGTGTGGCAGCAAAGAAATGAAAGGAAGTGTCTGGCAGCGCAGG
GGAGAAATTGTGTTTCTCCTTCTCCAAAGGTTACCTTTTCTATTTACCTTCATCATTATAAGAAGGTGAAA
GGATGCCAGGAGCAAGGCGGATCTACGGGACTTTTGAAAAGAAATGTACTGACCTTTGGCTGTAAAACTGTGTC
ATGAGAACAGAAAGCCCTGAGAGAAGAAGACAGCCATTATGGGATCTATGGCGCACTCTGTGCTAGGGCAGAAT
GCCAGGGAAGAGCTGCCTGTCTTCAGCGCCCTTGCTGGGAACCACACGCCAGAACCCGACCCTGCCATGCTTTGT
GCTTCAGGCAGCAGGACCCACCCCTAACAGAGTCTCAAGTCAGACACTAA

670/6881
FIGURE 623

MAGTKMGREDSVLDTAWGGSAAARETHAGAALSLVSLPSTGLRCGGTTERTSLRVSSALDGVVEEPWKLYLQKLTA
ALSLAAGKWERESKRKGGKGGSPCGQQERVHDSRIKTKALWSEFRKAEGDTKGGDKAQVKDEPQNR SARLSAKP
APPKPEPKPKKAPAETVPKGGKGA VAGKEGNNPAENGDAKTDQAQKAEGAGDAKANLLPRCGSKEMKGSVWQRR
GEIVFPSPKVLHLYFTFI IHKKVKGCQEQQGGSTGLLKRNVLTFGCKNCVMRTESFERRRQPFMGSMHSLV LGQN
AREELPVFSALAGNHTPEPDPAMLCASGSRTPLTESQVRH

671/6881
FIGURE 624

AAGAAGCTGCCGTTGTTCTGGGTACTACAGCAGAAGGGTATGCGGAAGCGAGCACCCAGTCTGAGATGGCTCCT
GCCGGTGTGAGCCTGAGGGCCACCATCCTCTGCCTCCTGGCCTGGGCTGGCCTGGCTGCAGGTGACCGGGTGTAC
ATACACCCCTTCCACCTCGTCATCCACAATGAGAGTACCTGTGAGCAGCTGGCAAAGGCCAATGCCGGGAAGCCC
AAAGACCCACCTTCATACCTGCTCCAATTCAGGCCAAGACATCCCCTGTGGATGAAAAGGCCCTACAGGACCAG
CTGGTGCTAGTCGCTGCAAACTTGACACCGAAGACAAGTTGAGGGCCGCAATGGTCGGGATGCTGGCCAACCTTC
TTGGGCTTCCGTATATATGGCATGCACAGTGAGCTATGGGGCGTGGTCCATGGGGCCACCGTCCTCTCCCAACG
GCTGTCTTTGGCACCTGGCCTCTCTCTATCTGGGAGCCTTGGACCACACAGCTGACAGGCTACAGGCAATCCTG
GGTGTTCCTTGAAGGACAAGAACTGCACCTCCCGGCTGGATGCGCACAAGGTCTGTCTGCCCTGCAGGCTGTA
CAGGGCTGCTAGTGGCCCAGGGCAGGGCTGATAGCCAGGCCAGCTGCTGTCTTCCACGGTGGTGGGCGTGTTC
ACAGCCCCAGGCCTGCACCTGAAGCAGCCGTTTGTGACGGGCTGGCTCTCTATACCCCTGTGGTCTCCACGC
TCTCTGGACTTCACAGAAGTGGATGTTGCTGCTGAGAAGATTGACAGGTTTCATGCAGGCTGTGACAGGATGGAAG
ACTGGCTGCTCCCTGATGGGAGCCAGTGTGGACAGCACCTGGCTTTCAACACCTACGTCCACTTCCAAGGGAAG
ATGAAGGGCTTCTCCCTGCTGGCCGAGCCCCAGGAGTTCTGGGTGGACAACAGCACCTCAGTGTCTGTTCCCATG
CTCTCTGGCATGGGCACCTTCCAGCACTGGAGTGACATCCAGGACAACCTTCTCGGTGACTCAAGTGCCTTCACT
GAGAGCGCTGCCTGCTGCTGATCCAGCCTCACTATGCCTCTGACCTGGACAAGGTGGAGGGTCTCACTTTCCAG
CAAACTCCCTCAACTGGATGAAGAACTGTCTCCCGGACCATCCACCTGACCATGCCCCAACTGGTGTGCAA
GGATCTTATGACCTGCAGGACCTGCTCGCCAGGCTGAGCTGCCCGCCATTCTGCACACCGAGCTGAACCTGCAA
AAATTGAGCAATGACCGCATCAGGGTGGGGGAGGTGCTGAACAGCATTTTTTTTGAGCTTGAAGCGGATGAGAGA
GAGCCACAGAGTCTACCCAACAGCTTAACAAGCCTGAGGTCTTGGAGGTGACCTGAACCGCCCATTCCTGTTT
GCTGTGTATGATCAAAGCGCCACTGCCCTGCACTTCTTGGCCGCGTGGCCAACCCGCTGAGCACAGCATGAGGC
CAGGGCCCCAGAACACAGTGCCTGGCAAGGCCTCTGCCCTGGCCTTTGAGGCAAAGGCCAGCAGCAGATAACAA
CCCCGGACAATCAGCGATGTGTACCCCCAGTCTCCACCTTTTCTTCTAATGAGTCGACTTTGAGCTGGAAAG
CAGCCGTTTCTCCTTGGICTAAGTGTGCTGTCATGGAGTGAGCAGTAGAAGCCTGCAGCGGCACAAATGCACCTCC
CAGTTTGCTGGGTTTATTTTAGAGAATGGGGTGGGGAGGCAAGAACCAGTGTTTAGCGGGGACTACTGTTCCA
AAAAGAATTCAAACCGACCAGCTTGTTTGTGAAACAAAAAAGTGTCCCTTTTCAAGTTGAGAACAAAAATTGGG
TTTTAAATTAAGTATACATTTTGCATTGCCTTCGGTTTGTATTAGTGTCTTGAATGTAAGAACATGACCTC
CGTGTAGTGTCTGTAATACCTTAGTTTTTCCACAGATGCTTGTGATTTTGAACAATACGTGAAAGATGCAAGC
ACCTGAATTTCTGTTTGAATGCGGAACAATAGCTGTTATTTCTCCCTTGTTAGTAATAAACGTCTTGCCAC

672/6881

FIGURE 625

MRKRAPQSEMAPAGVSLRATILCLLAWAGLAAGDRVYIHPFHLVIHNESTCEQLAKANAGKPKDPTFIPAPIQAK
TSPVDEKALQDQLVLVAAKLDTEKLRAMVGMLANFLGFRIYGMHSELWGVVHGATVLSPTAVFGTLASLYLGA
LDHTADRLQAILGVPWKDKNCTSRDAHKVLSALQAVQGLLVAQGRADSQAQLLLSTVVGVTAPGLHLKQPFVQ
GLALYTPVVLPRSLDFTELDVAAEKIDRFMQAVTGWKTGCSLMGASVDSTLAFNTYVHFQGKMKGFSLLAEPQEF
WVDNSTSVSPMLSGMGTFQHWSDIQDNFSVTQVPFTESACLLLIQPHYASDLKVEGLTFQNSLNWMKKLSPR
TIHLTMPQLVLQGSYDLQDLQAELPAILHTELNLOKLSNDRIRVGEVLNSIFFELEADEREPTTESTQQLNKPE
VLEVTILNRPFLLFAVYDQSATALHFLGRVANPLSTA

673/6881
FIGURE 626

GAATTCGGCAGGACCGGGATCCTGTGTAGCGGCTGCAGAGGGTGCCGCCGCCCTAGGCGAAGTAGGGCCGTCCT
GAGCGAAAGAACCGCCCCCAGCAGGAGCACCACCACGGCTTAGCAAAGAATCCCAGACCCCGCCCGGAAGGCAG
CCGCACCATGGAGTCTTCCAGTTTCACTCTTATTCTCCGTTGGCCCAACCAGTCCCAGCGCTGTCGTGCT
CCTCTACTCGAAGGAGCTCAAAAAGTGGGATGAGTTTGAAGATATTTAGAAAGAGAGGAGGCATGTCAGTGACTT
GAAATTTGCAATGAAATGCTACACACCTCTTGTCTATAAGGGAATTACTCCATGTAAACCAATTGATATTAAATG
TAGTGTTCTCAATTCTGAGGAGATTCAATTATGTCATTAAACAGCTTTCCAAGGAATCCCTTCAATCTGTGGATGT
CCTCCGAGAGGAAGTGAGTGAGATCTTAGATGAAATGAGTCACAACTGCGTCTTGGAGCCATTCGGTTTTGTGC
CTTCACCCTGAGCAAAGTATTTAAACAAATTTTCTCGAAGGTGTGTGTAATGAAGAAGGTATTGAGAACTACA
AAGAGCCATCCAGGAGCATCCTGTTGTTCTGCTGCCTAGTCATCGAAGTTACATTGACTTCCTCATGTGTGCTTT
TCTTCTATACAATTATGATTTGCTGTGCCAGTTATAGCAGCAGGAATGGACTTCCTGGGAATGAAATGGTTGG
TGAGCTGCTACGAATGTCGGGTGCCTTTTTCATGCGGCGTACCTTTGGTGGCAATAAACTCTACTGGGCTGTATT
CTCTGAATATGTAAAACTATGTTACGGAATGGTTATGCTCCTGTTGAATTTTCTCGAAGGGACAAGAAGCCG
CTCTGCCAAGACATTGACTCCTAAATTTGGTCTTCTGAATATTGTGATGGAGCCATTTTAAAAAGAGAAGTTT
TGATACCTACCTTGTCCCAATTAGTATCAGTTATGATAAGATCTTGAAGAACTCTTATGTGTATGAGCTTCT
AGGGGTTCTTAAACAAAAGAGTCTACAACCTGGGTGCTGAAAGCCAGAAAGATTCTCTGAAAATTTTGAAG
CATCCATGTGTACTTTGGAGATCCTGTGTCACTTCGATCTTTGGCAGCTGGGAGGATGAGTCGGAGCTCATATA
CTTGGTTCCAAGATACATTCTCAGAAACAGTCTGAGGACATGCATGCCTTTGTCACTGAAGTTGCCTACAAAAT
GGAGCTTCTGCAAATTGAAAACATGGTTTTGAGCCCTGGACCCTAATAGTTGCTGTTCTGCTTCAGAACCGGCC
ATCCATGGACTTTGATGCTCTGGTGAAAAGACTTTATGGCTAAAAGGCTTAACCCAGGCATTTGGAGGGTTCT
CATTTGGCCTGATAATAAACCTGCTGAAGAAGTTGTCCCGGCCAGCATTCTTCTGCATTCCAACATTGCCAGCCT
TGTCAAAGACCAGGTGATTCTGAAAGTGGACTCCGGAGACTCGGAAGTGGTCGATGGGCTTATGCTCCAGCACAT
CACTCTCCTCATGTGCTCAGCTTATAGGAACAGCTGCTCAACATTTTGTGCGCCCATCCTTAGTAGCAGTAGC
ATTGCAGATGACACCAGGGTTCAGGAAAGAGGATGTCTACAGTTGCTTTCGCTTCTACGTGATTTTTTGAGA
TGAGTTTCACTTCTTCCAGGAAACACACTAAAGGACTTTGAAGAAGGCTGTACCTGCTTTGTAAGTGAAGC
CATACAAGTGACTACGAAAGACATCCTAGTTACAGAGAAAGGAAATACTGTGTTAGAAATTTTAGTAGGACTCTT
TAAACCTTTTGTGAAAGCTATCAGATAATTTGCAAGTACCTTTTGAAGTGAAGAAGAGGACCCTTCACTGAGGA
ACAGTACTTGGCTGCAGTCAGAAAATTCACAAGTCAGCTTCTCGATCAAGGTACCTCTCAATGTTATGATGATT
ATCTTCTGATGTGCAGAAAAACGCCTTAGCAGCCTGTGTGAGGCTCGGAGTAGTGAGAAGAAGAAGATAAATAA
TAACTGTATATTTAATGTGAATGAACCTGCCACAACCAAATTAGAAGAAATGCTTGGTTGTAAGACACCAATAGG
AAAACCAGCCACTGCAAACTTTTAATAATCAACAAATAGTTATGGAATAATTCGGTCACGTAATTACTCTCATCGA
AGGACTCATTACAACAAACAGGGAAGTAAAGGAAGAGACACATCCTCTCATACTCCCTGAGACTCTGAGAACAGT
GGACGCAGAGGGAAGAGATGATCATTGGAAGCAATCAGTTTACTCTTCCCCACCACAGTGGTTAAAGGCGTTTG
TATCTGACACTATGTGTGTGTTTTAAATAAACTTTTGGAAACATGAAAAAAAAAAAAAAAAAACTCGAG

674/6881

FIGURE 627

MESSSSSNSYFSVGPTSPSAVVLLYSKELKKWDEFEDILEERRHVSDLKFAMKCYTPLVYKGITPCKPIDIKCSV
LNSEEIHYVIKQLSKESLQSVDLREEVSEILDEMSHLRLGAIRFCAFTLSKVFKQIFSKVCVNEEGIQKLQRA
IQEHPVLLPSHRSYIDFLMLSFLLYNYDLPVPVIAAGMDFLGKMGVCELLRMSGAFFMRRTFGGNKLYWAVFSE
YVKTMLRNGYAPVEFFLEGTRSRSAKTLTPKFGLLNIVMEPFFKREVFDTYLPISISYDKILEETLYVYELLGV
PKPKESTTGLLKARKILSENFGSIHVYFGDPVSLRSLAAGRMSRSSYNLVPRYIPQKQSEDMHAFVTEVAYKMEL
LQIENMVLSPWTLIVAVLLQNRPSMDFDALVEKTLWLKGLTQAFGGFLIWPDNKPAEEVVPASILLHSNIASLVK
DQVILKVDSGDSEVVDGLMLQHITLLMCSAYRNQLLNIFVRPSLVAVALQMPGFRKEDVYSCFRFLRDVFADEF
IFLPGNTLKDFEEGCYLLCKSEAIQVTTKDILVTEKGNTVLEFLVGLFKPFVESYQIICKYLLSEEDHFSEEQY
LAAVRKFTSQLLDQGTSCQYDVLSSDVQKNALAACVRLGVVEKKKINNNCIFNVNEPATTKLEEMLGCKTPIGKP
ATAKL

675/6881

FIGURE 628

CTCTTCCTAAGCCGGCGCTCGGCAAGTTCTCCAGGAGAAGGCCATGTCAGTACGAGCGCCAAGATCGTGAAGC
CTAATGGCGAGAAGCCGGACGAGTTCGAGTCCCGCATCTCCAGGCTCTTCTGGAGCTGGAGATGAACTTGGACC
TCAAGGCTCAGCTCAGGGAGCTGAATATTATGGCAGCCAAGGAAATTGAGGTTGGTGGTGGTCAGAAAGCTATCA
TAATCTTTGTTCCCGTTCTCAACTGAAATCTTTCCAGAAAATCCAAGTCCGGCTAGTATGTGAATTGGAGAAAA
AGTTCAGTGGGAAGCATGTTGTCTTTATCGCTTAAAGGAGAATCCTGCCTAAGCCAACTTGGAAAAGCTGTACAA
AATATAAGCAAAAGCATCCAGGAGCCATACTCTGACAGCTGTGCACGATGCCATCCTTGAGGACTTGGTCTTCA
CAAGCAAAATTGTGGGCAAGAGAATCCGCGTGAAACTAGATGGCAGCCGGCTCATGAAGGTTCAATTGGACAAAG
CACAGCAGAACAATGTGGAACACAAGGTTGAAACTTTTTCTGGTGTCTATAAGAAGCTCATGGGCAAGGATGTTA
ATTTGAATTCCAGAGTTTCAATTGTAACAAAAATGACTAAATAAAAAATATATATTACAGT

676/6881
FIGURE 629

MFSTSAKIVKPNGEKPDEFESRISQALLELEMNLDLKAQLRELNIMAAKEIEVGGGQKAIIFVPVPQLKSFQKI
QVRLVCELEKKFSGKHVVFIAKRRILPKPTWKSTKYKQKHRSHTLTAVHDAILEDLVFTSKIVGKRIRVKLDG
SRLMKVHLDKAQQNNVEHKVETFSGVYKKLMGKDVNFEPFQQL

677/6881
FIGURE 630

GTCGCGACCCTGGTCCGGACCTGACCTGAATTGCGACCCCAACCTGGACTGCTCCCCTGACCGCAACCCCTACCC
CCGCCCACCAGTATGGCCCCGGCACGTGTTCCCTAACGGGGCCCCCAGGAGTTGGAAAAACAACATTGATCCATAAA
GCCAGTGAGGTTTTAAATCCTCTGGTGTGCCTGTTGATGGATTTTATACCGAAGAAGTCAGACAGGGAGGGAGA
AGAATAGGATTTCGATGTCGTCACGTTGTCCGGCACCCGGGGGCCCTTTATCGAGAGTTGGGTTAGAGCCTCCACCT
GGAAAACGTGAATGCCGAGTTGGGCAGTATGTGGTCGACCTGACTTCTTTTGAGCAGTTGGCACTACCCGCTCTTG
AGGAATGCCGACTGCAGCAGTGGCCCAGGGCAAAGAGTGTGCGTCATCGATGAGATTGGGAAGATGGAGCTCTTC
AGTCAGCTTTTCATTCAAGCTGTTTCGTCAGACGCTGTCTACCCCAGGGACTATAATCCTTGGCACAATCCCAGTT
CCTAAAGGAAAGCCACTGGCTCTTGTAGAAGAAATCAGAAACAGAAAGGATGTGAAGGTGTTAATGTCACCAAG
GAAAACAGAAACCACCTTCTGCCAGATATCGTGACGTGCGTGCAGAGCAGCAGGAAGTGAAGACACGTGCATTCC
TGCCTTCCGTGAAGGAGTGCCCAGTTCAAGAGGAGCCTGATGGAGCCCTGCCTGTCGAGGCTGTATGCCCTATGGG
GTTATGGAACCTTGTGGGCTTTTCTAGAGAAACTCAACAGCTGTTTCCCATAAATGTTTAAAGATCAAATTA
GCCTTAATGCTGGATTGTCTGTACAAGATTAACTATCCATTGTGGCTTATCTATGCTTAAAGATTTCTTGTTAT
TAAAAAAAAAAAAAAAAAAAAAAAAA

678/6881

FIGURE 631

MARHVFLTGPPGVGKTTLIHKASEVLKSSGVPVDGFYTEEVQRQGGRRIGFDVVTLSGTRGPLSRVGLEPPPGKRE
CRVGQYVVDLTSFEQLALPVLARNADCSSGPGQRVCVIDEIGKMELFSQLFIQAVRQTLSTPGTIILGTIPVPKGK
PLALVEEIRNRKDVKFNVTKENRNHLLPDIVTCVQSSRK

679/6881
FIGURE 632

GCAAGATGGCAGAAGTAGAGCAGAAGAAGAAGCGGACCTTCCGCAAGTTCACCTACCGCGGCGTGGACCTGGACC
AGCTGCTGGACATGTCTACGAACAGCTGATGCAGCTGTACAGTGCGCGCCAGCGGCGGCGGCTGAACCGGGGCC
TGCGGCGGAAGCGGCACTACCTGCTGAAGCGCCTGCGCAAGGCCAAGAAGGAGGCGCGCCCATGGAGAAGCCGG
AAGTGGTGAAGACGCACCTGAGGGACGTGATCATCCTGCCCCGAGGTGGTGGGCAGCATGGTGGGCGTCCACAACG
GCAAGACCTTCACCTACAAGCCCGTAAAGCAATGCCGGTCCGGCATCGGCGCCACCCACTCCTTCTGCTTCATCC
CTCTCAAGTAGTGGCTCAGCTAATAAAGGCGCACATG

680/6881
FIGURE 633

MAEVEQKKKRTFRKFTYRGVDLDQLLDMSYEQMLQLYSARQRRRLNRGLRRKRHYLLKRLRKAKKEAPPMEKPEV
VKTHLRDVIIILPEVVGSMVGVHNGKTFTYKPVKQCRSGIGATHSFCFIPLK

681/6881

FIGURE 634

ACCAATGTCTATTCTCAAGATCCATGCCAGGGAGCTCTTTGACTCTCGTGGGAATCCCACTGTTGAGGTTGATCTC
TTCACCTCAGAAGGTCTCTTCAGAGCTGCTGTGCCCAGTGGTGCTTCAACTGGTATCTATGAGGTCCTAGAGCTC
CAGGACAATGATAAGACTCGCTATATGGGGAAGGGTGTCTCAAAGCCTGTTGAGCCCATCAATAAACTATTGCA
CCTGTCCTGGTTAGCAAGAACTGAACGTACAGAACTTCTACAGAAGCCAAGCTCCCTGGAGCCCTGTTGGCAG
CTCTAGCCTTGCAGTCATGTAATTGGCCCAAATCACCGGAGCCACGTGACCCTCCAGTGTCATCTCCGGGGTGGC
CACAGGCAAGATCCCCAGTGATTTTGTGCTCAAAATAAAAAGCCTCATTGACCCATGAG

682/6881
FIGURE 635

MSILKIHARELFDNRGNPTVEVDLFTSEGLFRAAVPSGASTGIYEVLELQDNDKTRYMGKGVSKPVEPINKTIAP
VLVSKKLNVTPELLQKPSSLEPCWQL

683/6881
FIGURE 636

CCCAGAAATTCTACCCAAGCTCCCTCAGCACCATGTACCGAGCACTTCGGCTCCTCGCGCGCTCGCGTCCCCTCG
TGC GGGCTCCAGCCG CAGCCTTAGCTTCGGCTCCCGGCTTGGGTGGCGCGGCCGTGCCCTCGTTTTGGCTCCGA
ACGCGGCTCGAATGGCAAGCCAAAATTCCTCCGGATAGAATATGATACCTTTGGTGAATAAAGGTGCCAAATG
ATAAGTATTATGGCGCCAGACCGTGAGATCTACGATGAACCTTAAGATTGGAGGTGTGACAGAACGCATGCCAA
CCCCAGTTATTAAAGCTTTTGGCATCTTGAAGCGAGCGGCCGTGAAGTAAACCAGGATTATGGTCTTGATCCAA
AGATTGCTAATGCAATAATGAAGGCAGCAGATGAGGTAGCTGAAGGTAAATTAATGATCATTTCCTCTCGTGG
TATGGCAGACTGGATCAGGAATCAGACAAATATGAATGTAAATGAAGTCATTAGCAATAGAGCAATTGAAATGT
TAGGAGGTGAACCTTGGCAGCAAGATACCTGTGCATCCCAACGATCATGTTAATAAAAGCCAGAGCTCAAATGATA
CTTTTCCCACAGCAATGCACATTGCTGCTGCAATAGAAGTTCATGAAGTACTGTTACCAGGACTACAGAAGTTAC
ATGATGCTCTTGATGCAAAATCCAAAGAGTTTGACAGATCATCAAGATTGGACGTACTCATACTCAGGATGCTG
TTCCACTTACTCTTGGGCAGGAATTTAGTGGTTATGTTCAACAAGTAAATATGCAATGACAAGAATAAAAGCTG
CCATGCCAAGAATCTATGAGCTCGCAGCTGGAGGCACTGCTGTTGGTACAGGTTTAAATACTAGAATTGGCTTG
CAGAAAAGGTTGCTGCAAAAGTGGCTGCACCTTACAGGCTTGCTTTTGTACTGCTCCGAATAAATTTGAAGCTC
TGGCTGCTCATGACGCTCTGGTTGAGCTCAGTGGAGCCATGAACACTACTGCCTGCAGTCTGATGAAGATAGCAA
ATGATATTGATTTTTGGGTCTCGGTCCGTGAGGTCTGGGAGAATTGATCTTGCTGAAAATGAACCAGGAA
GCAGTATCATGCCAGGCAAGGTGAACCTACTCAGTGTGAAGCAATGACCATGGTTGCAGCCCAAGTCATGGGGA
ACCATGTTGCTGTCACTGTGCGAGGCAGCAATGGACATTTTGAGTTGAATGTTTTCAAGCCAATGATGATTAAAA
ATGTGTTACACTCAGCCAGGCTGCTGGGGGATGCTTCAGTTTCCTTTACAGAAAAGTGCCTGGTGGGAATCCAGG
CCAATACAGAAAGGATCAACAAGCTGATGAATGAGTCTCTAATGTTGGTGACAGCTCTCAATCCTCATATAGGGT
ATGACAAGGCAGCAAAGATTGCTAAGACAGCACAAAAATGGATCAACCTTAAAGGAAAGTCTATCGAACTTG
GCTATCTCACAGCAGAGCAGTTTGACGAATGGGTAAACCTAAGGACATGCTGGGTCCAAAGTGAATTACATAAA
TTTATAATGAAAATAAACATGTATAAAATTTAAAAAACAGACTCCCATTTCTTAAAAACGGATAAGTTTGAAG
GAAACTGCTATTGAACCTAAGCATCTCTAGCAGAGCAATTTGATCAGTATATAAAACCCTAGGATGTGCTAGGTC
TAAGATGGATTAAACAAGTATAAAATAAAATACATTTATAAAATAAAAGGAAAACAGACTTAAAA

684/6881

FIGURE 637

MYRALRLLARSRLVRAPAAALASAPGLGGAAPFSFWPPNAARMASQNSFRIEYDTFGELKVPNDKYYGAQTVRS
TMNFKIGGVTERMPTPVIKAFGILKRAAAEVNQDYGLDPKIANAIMKAADEVAEGKLNDFPLVVWQTGSGTQTN
MNVNEVISNRAIEMLGELGSKIPVHPNDHVNKSQSSNDTFPTAMHIAAAIEVHEVLLPGLQKLHDALDAKSKEF
AQIIKIGRTHQTQDAVPLTLGQEFSGYVQQVKYAMTRIKAAMPRIYELAAGGTAVGTGLNTRIGFAEKVAAKVAAL
TGLPFVTAPNKFEEALAAHDALVELSGAMNTTACSLMKIANDIRFLGSGPRSGLGELILPENEPGSSIMPGKVNPT
QCEAMTMVAAQVMGNHVAVTVGGSNGHFELNVFKPMMIKNVLHSARLLGDASVSFTENCVVGIQANTERINKLMN
ESLMLVTALNPHIGYDKAAKIAKTAHKNGSTLKETAIELGYLTAEQFDEWVKPKDMLGPK

685/6881
FIGURE 638

GGGXTCGGCCCCCTCCCCCTCTTCTGCGCTCTCTTCGGGATACACGTGGGCTTCGGGCCTGGGCCGCGCAGTITTTT
CTTTGGCTTTCTCGAATCATCGAGGAGACTGGCAAAATTAGTAATGTCTTTAGACTGCGTCACTAGCGCCAATCG
AATTCGTAGTTGATTGTTCTTTGAATTCACATTGAGTCCTCGGTTAAAACCTCAGTTAAAAAGCAGGATAAAGT
CGAGCTGCTTTGGTTCTCGGAACGGAAAAGCGTTTTTTTTGTTGTTTAGAGGCTTGGCCTTTATGGGGCTCGTG
TGTTTTAGGGGACGGCAAAACAGAACAGAAAGCGGAGATAAAAAGAGGGGTGTTAAAAGACCACGAGAAGATCA
TGGCCGTGGATATTTTGAAGTACATTGAAGAGAACAAAGTATAGCAGAGCCAAATCTCCTCAGCCACCTGTTGAAGA
AGAAGATGAACACITTCGATGACACAGTGGTTTGTCTTGATACTTATAATTGTGATCTACATTTTAAAAATATCAAG
AGATCGTCTCAGTGCTTCTTCCCTTACAATGGAGAGTTTTGCTTTTCTTTGGGCTGGAGGAAGAGCATCCTATGG
TGTGTCAAAAGGCAAAAGTGTGTTTTGAGATGAAGGTTACAGAGAAGATCCCAGTAAGGCATTTATATACAAAAGA
TATTGACATACATGAAGTTCGTATTGGCTGGTCACTAACTACAAGTGAATGTTACTTGGTGAAGAAGAATTTTC
TTATGGGTATTCTCTAAAAGGAATAAAAACATGCAACTGTGAGACTGAAGATTATGGAGAAAAGTTTGATGAAAA
TGATGTGATTACATGTTTTGCTAACTTTGAAAGTGATGAAGTAGAACTCTCGTATGCTAAGAATGGACAAGATCT
TGGCGTTGCGCTTCAAAATCAGTAAGGAAGTCTTGCTGGACGGCCACTGTTCCCGCATGTTCTCTGCCACAACCTG
TGCAGTTGAATTTAATTTTGGTCAGAAGGAAAAGCCATATTTTCCAATACCTGAAGAGTATACTTTTCATCCAGAA
CGTCCCCCTTAGAGGATCGAGTTAGAGGACCAAAGGGGCTGAAGAGAAGAAAGATTGTGAAGTTGTGATGATGAT
TGGCTTGCCAGGAGCTGGAAAACTACCTGGGTACTAAACATGCAGCAGAAAATCCAGGGAAATATAACATTCT
TGGCACAATACTATTATGGATAAGATGATGGTGGCAGGTTTTAAGAAGCAAATGGCAGATACTGGAAAACTGAA
CACACTGTTGCAGAGAGCCCCCAGTGCTTGGGAAATTTATTGAGATTGCTGCCGAAAGAAGCGAAATTTTAT
TCTGGATCAGACAAATGTGTCTGCTGCTGCCAGAGGAGAAAAATGTGCTGTTTGCAGGCTTCCAGCGAAAAGC
TGTTGTAGTTTGGCCAAAAGATGAAGACTATAAGCAAAGAACACAGAAGAAAGCAGAAGTAGAGGGGAAAGACCT
ACCAGAACATGCGGTCCTCAAAATGAAAGGAACTTTACCCTCCCAGAGGTAGCTGAGTGCTTTGATGAAATAAC
CTATGTTGAACTTCAGAAGGAAGAAGCCCCAAAACCTCTGGAGCAATATAAGGAAGAAAGCAAAAAGGCTCTTCC
ACCAGAAAAGAAACAGAACACTGGCTCAAAGAAAAGCAATAAAAATAAGAGTGGCAAGAACCAGTTTAAACAGAGG
TGGTGGCCATAGAGGACGTGGAGGATTCAATATGCGTGGTGGAAATTTAGAGGAGGAGCCCCCTGGGAATCGTGG
CGGATATAATAGGAGGGGCAACATGCCACAGAGAGGTGGTGGCGGTGGAGGAAGTGGTGAATCGGCTATCCATA
CCCTCGTGCCCTGTTTTTCTGGCCGTGGTAGTTACTCAAACAGAGGGAAGTACAACAGAGGTGGAATGCCCAA
CAGAGGGAAGTACAACAGAACTTCAGAGGACGAGGAAACAATCGTGGCTACAAAATCAATCTCAGGGCTACAA
CCAGTGGCAGCAGGGTCAATCTGGGGTCAGAAGCCATGGAGTCAGCATTATCACCAAGGATATTATTGAATACC
CAAATAAAACGAAGTATACATATTTCTCCAAAACCTTACAAGAAGTCGACTGTTTTCTTTAGTAGGCTAACTT
TTTAAACATTCCACAAGAGGAAGTGCCTGCGGTTCTTTTTTAGAAGCTTTGTGGGTGATTTTTTTCTTTTC
TTTTTTGTACATTTTAAATTGCAGTTTAAAAGTGAATCGTAAGAGAACCTCAGCATTGTGCACGATAAGAGAATG
TGTCAGTATTTAGGGTTCTACATTTTATCTGTAAATGTGACTTTTTTTTTTTTTTATCACAAACAGAAGTAAAA
TGTTGCTTTGTACCTGGTGTCTTTTATTAAGAATTTACTCCCCCATTTCTCACAGAGAATAACAGTCGGGAGTC
ATTGTCACAATATAATAGAAATGTTAGCAACCAGATTGATGAAGGACTAAGTGGTCCTCATGAATTGCATTAG
ACTCTGTACTGCTCATATTACACTCCATCCTCTCTGTAGTTTGGTGGTAGTGGAGGGGTAAAGCTAAATCATAG
TTTCTGACAATAACTGGGAAGGTTTTTCTTAAATAACAATGGAATTGGTATAATTGGGATTGAAAACATAAAC
TTGGAACATAAGATAGAGAAGATGGAGTGTATGTAGAAGGCTGTTAAAAATGTAAACTTGGTTGCATTATTTGT
GGAGGCTCAAACCTTGTGAAGGTTAATACCATAATTTTCCATTGTTCTGCATTTTGATTCTGAAAAGAAAGCTG
GCTTTGCCCATTCTTATTAAAAAACTGTTGTAAATCCAGTTGTCTAATGGGATCTATATGAAGTTAGCCATG
TCTGTATGCCCTTCTCCACAAAATACTGTATAACTAGTGTGCTTGTAGTAGTTAACTCCACCATCTTTGTAAGC
TAATGAAATTGTGAGTCACCCATTTATATCTTAATTTTAAATCATGTGAGTTCTTGAATGGGTATCTCCTTAGCC
TGCTGATTTCTTTTTCTTTCTAAAGAAAGTGGGTGGAGAAATTAATTTAGACGTTTGTGTTGCAATAAAAAGAATT
CATTTT

686/6881
FIGURE 639

GGCGTTGCCGGCCGTGGGTGCTCTGGCCACAGTGAGTTAGGGGCGTCGGAGCGGGTTTCTCCAACCGCAATCGGC
TCCGCTCAAGGGGAGGAGAGAGTCCCTTCTCGGAAGGCCTAAGGAAACGTGTCGTCTGGAATGGGCTTGGGGGC
CACGCCTGCACATCTCCGCGAGACAGAGGGATAAAGTGAAGATGGTGCTGTTATTGTTACCTCGAGTGCCACATG
CGACCTCTGAGATATGTACACAGTCATTCTTACTATCGCACTCAGCCATTCTTACTACGCTAAAGAAGAAATAAT
TATTCGAGGATATTGCTGGCCGAGAGAACTTATGTAAATTTTCATGAAGTATTATATCCGTTTCTCTGGAG
TGAGAGAAACTCTTTTTAGATATCATCTGAGAGAACTAGTGAATCCAGTCACTGAGTGGAGTTGAGAGTCTAA
GAACCTCTGAAATTTGAGAACTGCTGGACCAGAGCCTTATAGAGCTCTGATAAGGTGTCAACAGGGTAGTTAATTT
GGCACCATGGGGGATACAGGGATTGCTACAATTTATCAAAGAAGCTTCAGAACCCATCCATGTGAGGAAGTATAAA
GGGCAGGTAGTAGCTGTGGATACATATTGCTGGCTTCACAAAGGAGCTATTGCTTGTGCTGAAAACTAGCCAAA
GGTGAACCTACTGATAGGTATGTAGGATTTTGTATGAAATTTGTAAATATGTTACTATCTCATGGGATCAAGCCT
ATTCTCGTATTGATGGATGTACTTTACCTTCTAAAAAGGAAGTAGAGAGATCTAGAAGAGAAAGACGACAAGCC
AATCTTCTTAAGGGAAAGCAACTTCTTCGTGAGGGGAAAGTCTCGGAAGCTCGAGAGTGTTCACCCGGTCTATC
AATATCACACATGCCATGGCCACAAAGTAATTAAGCTGCCCGGTCTCAGGGGGTAGATTGCCTCGTGGCTCCC
TATGAAGCTGATGCGCAGTTGGCCTATCTTAACAAAGCGGGAATTGTGCAAGCCATAATTACAGAGGACTCGGAT
CTCCTAGCTTTTGGCTGTAAAAAGGTAATTTTAAAGATGGACCAGTTTGGAAATGGACTTGAAATGATCAAGCT
CGGCTAGGAATGTGCAGACAGCTTGGGGATGTATTACGGAAGAGAAGTTTCGTTACATGTGTATTCTTTTCAGGT
TGTGACTACCTGTCTACTGCGTGGGATTGGATTAGCAAAGGCATGCAAAGTCCCTAAGACTAGCCAATAATCCA
GATATAGTAAAGTTATCAAGAAAATTGGACATTATCTCAAGATGAATATCACGGTACCAGAGGATTACATCAAC
GGGTTTATTTCGGGCCAACAACTTCTCTATCAGCTAGTTTTTGATCCCATCAAAGGAACTTATTCCTCTG
AACGCCATGAAGATGATGTTGATCTGAAACACTAAGCTACGCTGGGCAATATGTTGATGATTCCATAGCTCTT
CAAATAGCACTTGGAAATAAAGATATAAATACTTTTGAACAGATCGATGACTACAATCCAGCACTGCTATGCCT
GCCCCATTCAAGAAGTCATAGTTGGGATGACAAAACATGTCAAAGTCAGCTAATGTTAGCAGCATTTGGCATAGG
AATTACTCTCCAGACCAGAGTCGGGTACTGTTTCAGATGCCCCACAATTGAAGGAAATCCAACTACTGTGGGA
GTGGAACGAGTGATTAGTACTAAAGGGTTAAATCTCCCAAGGAAATCATCCATTGTGAAAAGACCAAGAAGTGCA
GAGCTGTGAGAAGATGACCTGTTGAGTCAGTATTCTCTTTCATTTACGAAGAAGACCAAGAAAAATAGCTCTGAA
GGCAATAAATCATTGAGCTTTTCTGAAGTGTGTTGTGCTGACCTGGTAAATGGACCTACTAACAAAAAGAGTGTA
AGCACTCCACCTAGGACGAGAAATAAATTTGCAACATTTTACAAAGGAAAAATGAAGAAAGTGGTGCAGTTGTG
GTTCCAGGGACCAGAAGCAGGTTTTTTTGCAGTTCAGATTCTACTGACTGTGTATCAAACAAAGTGAGCATCCAG
CCTCTGGATGAACTGCTGTACAGATAAAGAGAACAATCTGCATGAATCAGAGTATGGAGACCAAGAAGGCAAG
AGACTGGTTGACACAGATGTAGCACGTAATTCAAGTGATGACATTCCGAATAATCATATTCCAGGTGATCATATT
CCAGACAAGGCAACAGTGTGTTACAGATGAAGAGTCCCTACTCTTTTGAGAGCAGCAAATTTACAAGGACCATTTC
CCACCCACTTTGGGAACACTAAGAAGTTGTTTGTAGTTGGTCTGGAGGTCTTGGAGATTTTTCAAGAACGCCGAGC
CCCTCTCAAGCACAGCATTCGAGCAGTTCCGAAGAAAGAGCGATTCCCCACCTCTTGCCTGAGAATAATATG
TCTGATGTGTCGAGTTAAAGAGCGAGGAGTCCAGTGACGATGAGTCTCATCCCTTACGAGAAGGGGCATGTTCT
TCACAGTCCCAGGAAAGTGGAGAATTCTCACTGCAGAGTTCAAATGCATCAAAGCTTTCTCAGTGTCTAGTAAG
GACTCTGATTTCAGAGGAATCTGATTGCAATATTAAGTTACTTGACAGTCAAAGTGACCAGACCTCCAAGCTATGT
TTATCTCATTCTCAAAAAAAGACACACCTCTAAGGAACAAGTTCTGGGCTATATAAGTCCAGTTCTGCAGAC
TCTCTTTCTACAACCAAGATCAAACCTCTAGGACCTGCCAGAGCCAGTGGGCTGAGCAAGAAGCCGGCAAGCATC
CAGAAGAGAAAGCATCATAATGCCGAGAACAGCCGGGGTTACAGATCAAACCTCAATGAGCTCTGGAAAACTTT
GGATTTAAAAAAGATTCTGAAAAGCTTCTCTCTGTGAAGAAACCCCTGTCCCAGTCAGAGATAACATCCAATA
ACTCCAGAAGCGGAAGAGGATATATTTAACAACCTGAATGTGGCCGTGTTCAAAGAGCAATATTCCAGTAAATG
CAGACTGTCTCAAAGCTTTTGCCTGCAAGAGAATCTGATCAATTTGAAGTCCCTGTTTGGGAATGAGGCACTTAT
CAGCATGAAGAATTTTTTCTCATTCTGTGCCATTTTAAAAATAGAATACATTTTGTATATTGACTTTAAAAAAA
AAAAA

687/6881
FIGURE 640

MGIQGLLQFIKEASEPIHVRKYKGQVVAVDTYCWLHKGAIACAELAKGEPTDRYVGFCMKFVNMLLSHGKIPIL
VFDGCTLP SKKEVERSRERRQANLLKGKQLLREGKVSEARECFTRSINITHAMAHKVIKAARSQGVDCLVAPYE
ADAQLAYLNKAGIVQAIITEDSDLLAFGCKKVILKMDQFGNGLEIDQARLGMCRLGDVFTEEFKFRYMCILSGCD
YLSSLRGIGLAKACKVLRANNPDIKVIKKIGHYLMKNITVPEDYINGFIRANNTFLYQLVFDPIKRKLIPINA
YEDDVDPETLSYAGQYVDDSIQIALGNKDINTFEQIDDYNDTAMPASRSWSDDKTCQKSANVSSIWHRNY
SPRPESGTVSDAPQLKENPSTVGVERVISTKGLNLPRKSSIVKRPRSAELSEDDLLSQYSLSFTKKTKKNSSEGN
KSLSFSEVFPDLVNGPTNKKSVSTPPRTRNKFATFLQRKNEESGAVVVPGTRSRFFCSDSDTCVSNKVSIOPL
DETAVIDKENNLHESYGDQEGKRLVDTDVARNSSDDIPNNHIPGDHIPDKATVFTDEESYSFESSKFTRTISPP
TLGTLRSCFSWSGGLGDFSRTPSPSPSTALQQFRRKSDSPTSLPENMSDVSQKSEESSDDESHPLREGACSSQ
SQESGEFSLQSSNASKLSQCSSKSDSEESDCNIKLLDSQSDQTSKLCLSHFSKKDTPLRNKVPGLYKSSSADSL
STTKIKPLGPARASGLSKKPASIQRKHHNAENKPGLIKLNELWKNFGFKKDEKLPPCKKPLSPVRDNIQLTP
EAEEDIFNKPECGRVQRAIFQ

688/6881

FIGURE 641

TCTAATACCTATTGAICTGTCACTTTCTCCCATCACGCTCAGGTGGGACCATCCAGTTGCAGGAAAACAAGCTTA
ACACGCCCCTGATTCTACATTATGGTGAGTTCTATAATTATTTTATTATATATTACAGTGTAAATGGAATA
AAGTGCCTAATAAATGCAAATGTGCTTACATCTTTTGGCCCAGCTCCTACCTCCTGGCAGCCTCTCCAGGCCCAG
AACTTTCTCCAGTCAGCCTCTACAGACCAAGCTCATGACTCACAATGGCCTATTTAGGCCCATAACCCTACGTCAC
GGCAGTCTCCGCAGATGAGGCTACTGCCTCACAAACAGCCTCCACAGGCACAGCTCCATCGTTACAATGGCCTCTT
TAGACCCAGCTCCTGCCTCCCAGCCTTCTCTCCAGGCCCTGAACTTTCTCAAGTCGACCTCACCAGGCCCAGCTC
ATGCTTCTTTGCAGCCTCTCCAGGCCCAGCTCCTGCATCTTGGTGGCCCCCTCCAGGCCCAGCCTCTGCCTCCCGT
CGGCCTCTACAGTCCCAACATCTGTCTCACAGCAGATTCTTCAGGCCCAGCATCTGCCTCACTTGGACCCTCCAG
ACCCAGATGGTGTCTCACTGTGGCATCCTCAGGTGAAGCTCCTGCCTTTCGGCAGCCTCTCCAGGCCCAGCTCCT
CCTGCCTCCCAGTGGCCTCTTTTCGGCCCAGCCCAGCTCATGCCTCCCGGCGGCCTTCCCAAGCCCCGCTTTTGAC
TTTCCACCGAAAGTCCCAGCCTCCTGCCTCCCGAAGGCCTGCACAGGCCCA

689/6881
FIGURE 642

GTCCACATGCAGAGGCTCGCTACAGAAGCTACAGAAGAAGATGGTTACATTTC AAGTGGGCAGGTGCCGTTTGCT
TATGAGCACCATCAGCCACATCAGCCACGGTTCCTTTGAATTTTACCTAATTGTCTCTAATTCTTAACAATTCTTT
GAGGTGATCTGCAACTCTTTCACCATCTGTAATGCGGAGATGCAGGAAGTTGGTGTGGCCTATATCCCAGTATC
TCTTTGCTCAATCAGCTGTGACCCCAACTGTTTCGATTGTGTTCAATGGGCCCCACCTCTTACTGCGAGCAGTC
CGAGACATCGAGGTGGGAGAGGAGCTCACCATCTGCTACCTGGATATGCTGATGACCAGTGAGGAGCGCCGGAAG
CAGCTGAGGGACCAGTACTGCTTTGAATGTGACTGTTTCCGTTGCCAAACCCAGGACAAGGATGCTGATATGCTA
ACTGGTGATGAGCAAGTATGGAAGGAAGTTCAAGAATCCCTGAAAAAATTGAAGAACTGAAGGCACACTGGAAG
TGGGAGCAGGTTCTGGCCATGTGCCAGGCAATCATAAGCAGCAATTCTGAACGGCTTCCCGATATCAACATCTAC
CAGCTGAAGGTGCTCGACTGCGCCATGGATGCCTGCATCAACCTCGGCCTGTTGGAGGAAGCCTTGTTCTATGGT
ACTCGGACCATGGAGCCATACAGGATTTTTTCCCAGGAAGCCATCCCGTCAGAGGGGTTCAAGTGATGAAAGTT
GGCAAACCTGCAGCTACATCAAGGCATGTTTCCCAAGCAATGAAGAATCTGAGACTGGCTTTTGATATTATGAGA
GTGACACATGGCAGAGAACACAGCCTGATTGAAGATTTGATTCTACTTTTAGAAGAATGCGACGCCAACATCAGA
GCATCCTAAGGGAACGCAGTCAGAGGGAAATACGGCGTGTGTCTTTGTTGAATGCCTTATTGAGGTACACACTC
TATGCTTTGTTAGCTGTGTGAACCTCTCCTATTGGAAATTCTGTTCCGTGTTTGTGTAGGTAAATAAAGGCAGAC
ATGGTTTGCAAACCACAAGAATCATTAGTTGTAGAGAAGCACGATTATAATAAATTCAAAACATTTGGTTGAGGA
TGCC

690/6881
FIGURE 643

CTTGCTTCGGACGCCGGATTTTGACGTGCTCTCGCGAGATTGGGTCTCTTCCTAAGCCGGCGCTCGGCAAGTTC
TCCCAGGAGAAAGCCATGTTCAGTTTCGAGCGCCAAGATCGTGAAGCCCAATGGCGAGAAGCCGGACGAGTTCGAG
TCCGGCATCTCCCAGGCTCTTCTGGAGCTGGAGATGAACTCGGACCTCAAGGCTCAGCTCAGGGAGCTGAATATT
ACGGCAGCTAAGGAAATTGAAGTTGGTGGTGGTCGGAAAGCTATCATAATCTTTGTTCCCGTTCCTCAACTGAAA
TCTTTCCAGAAAATCCAAGTCCGGCTAGTACGCGAATTGGAGAAAAAGTTCAGTGGGAAGCATGTCGTCTTTATC
GCTCAGAGGAGAATTCTGCCTAAGCCAACCTCGAAAAAGCCGTACAAAAATAAGCAAAAGCGTCCCAGGAGCCGT
ACTCTGACAGCTGTGCACGATGCCATCCTTGAGGACTTGGTCTTCCCAAGCGAAATTGTGGGCAAGAGAATCCGC
GTCAAAC TAGATGGCAGCCGGCTCATAAAGGTTCAATTTGGACAAAGCACAGCAGAACAATGTGGAACACAAGGTT
GAAACTTTTTCTGGTGTCTATAAGAAGCTCACGGGCAAGGATGTTAATTTGAATTCCCAGAGTTTCAATTGTAA
ACAAAAATGACTAAATAAAAAGTATATATTCACAGTAAAAAAAAAAAAAAAAAAAA

691/6881
FIGURE 644

MFSSSAKIVKPNGEKPDEFESGISQALLELEMNSDLKAQLRELNITAAKEIEVGGGRKAIIFVVPVQLKSFQKI
QVRLVRELEKKFSGKHVVFIQRRILPKPTRKSRTKNKQKRPRSRTLTA VHDAILEDLVFPSEIVGKRIRVKLDG
SRLIKVHLDKAQQNNVEHKVETFSGVYKKLTGKDVNFEFPEFQL

692/6881
FIGURE 645

GGCACGAGGCGGGCCAGCGACGGGCAGGACGCCCCGTTTCGCCTAGCGCGTGCTCAGGAGTTGGTGTCTGCCTGC
GCTCAGGATGAGGGGGAATCTGGCCCTGGTGGGCGTTCTAATCAGCCTGGCCTTCCTGTCACTGCTGCCATCTGG
ACATCCTCAGCCGGCTGGCGATGACGCCTGCTCTGTGCAGATCCTCGTCCCTGGCCTCAAAGGGGATGCGGGAGA
GAAGGGAGACAAAGGCGCCCCGGACGGCCTGGAAGAGTCGGCCCCACGGGAGAAAAAGGAGACATGGGGGACAA
AGGACAGAAAGGCAGTGTGGGTCGTTCATGGAAAAATTGGTCCCATTGGCTCTAAAGGTGAGAAAGGAGATTCCGG
TGACATAGGACCCCCCTGGTCCTAATGGAGAACCAGGCCTCCCATGTGAGTGCAGCCAGCTGCGCAAGGCCATCGG
GGAGATGGACAACCAGGTCTCTCAGCTGACCAGCGAGCTCAAGTTCATCAAGAATGCTGTGCGCCGGTGTGCGCGA
GACGGAGAGCAAGATCTACCTGCTGGTGAAGGAGGAGAAGCGCTACGCGGACGCCAGCTGTCTGCCAGGGCCG
CGGGGGCACGCTGAGCATGCCCAAGGACGAGGCTGCCAATGGCCTGATGGCCGCATACCTGGCGCAAGCCGGCCT
GGCCCGTGTCTTCAICGGCATCAACGACCTGGAGAAGGAGGGCGCCTTCGTGTACTCTGACCACTCCCCATGCG
GACCTTCAACAAGTGGCGCAGCGGTGAGCCCAACAATGCCTACGACGAGGAGGACTGCGTGGAGATGGTGGCCTC
GGGCGGCTGGAACGACGTGGCCTGCCACACCACCATGTACTTCATGTGTGAGTTTGACAAGGAGAACATGTGAGSC
CTCAGGCTGGGGCTGCCCATTGGGGGCCCCACATGTCCCTGCAGGGTTGGCAGGGACAGAGCCCAGACCATGGTG
CCAGCCAGGGAGCTGTCCCTCTGTGAAGGGTGGAGGCTCACTGAGTAGAGGGCTGTTGTCTAAACTGAGAAAATG
GCCTATGCTTAAGAGGAAAATGAAAGTGTTCTGGGGTGCTGTCTCTGAAGAAGCAGAGTTTCATTACCTGTATT
GTAGCCCCAATGTCATTATGTAATTATTACCCAGAATTGCTCTTCCATAAAGCTTGTGCCTTTGTCCAAGCTATA
CAATAAAATCTTTAAGTAGTGCAGTAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA

693/6881

FIGURE 646

MRGNLALVGVLI SLAFLSLLPSGHPQPAGDDACSVQILVPGLKGDAGEKGDKGAPGRPGRVGPTGEKGDMDKGQ
KGSVGRHGKIGPIGSKGEKGDSDIGPPGPNGEPGLPCECSQLRKAIGEMDNQVSQLTSELKFIKNAVAGVRETE
SKIYLLVKEEKRYADAQLSCQGRGGTLSMPKDEAANGLMAAYLAQAGLARVFIGINDLEKEGAFVYSDHSPMRTF
NKWRSGEFNAYDEEDCVEMVASGGWNDVACHTTMYFMCEFDKENM

694/6881
FIGURE 647

GGCAGAGGGGCTTCTGTACCTTCCTCACGGACCTTGGTCACGGCCGCAGGTGACCCCTTAGCCCAGCTCCAGT
GGGCGGGTGGCAGGGTCATGGAGGACGCTGGCGGGCGCGAGGAGACCCCGGCCCCGAGGCCCCGCACCCCCCTC
AGCTCGGCCTCCGGAGGAGCAGGGGTTGCTCTTCCAGGAGGAAACCATCGATCTTGGCGGAGATGAGTTTGGAT
CCGAAGAGAACGAGACCGCATCGGAAGGCTCGAGTCCTCTCGCGGACAAGCTGAACGAACACATGATGGAGAGCG
TCCTCATCTCTGACTCCCCAACAGCGAGGGCGACGCGGGCGACCTGGGCGGAGTGCGGGACGAAGCTGAGCCCCG
GAGGGGAAGGCGACCCAGGCCCCGAGCCCCGCGGGCACCCCGAGTCCCAGCGGCGAGGCCGACGGCGACTGTGCCC
CCGAGGACGCGGCACCCAGTAGCGGAGGGGCCCCGAGGCAGGACGCGGCCCGCGAGGTCCCAGGCAGCGAAGCCG
CGCGCCCGGAGCAGGAGCCTCCCGTTGCGGAGCCGGTCCCGGTGTGCACCATCTTCAGCCAGCGCGCGCCCCCAG
CCTCCGGGGACGGCTTCGAGCCGCAGATGGTGAAGTCGCCAGCTTCGGTGGCGCCAGCGAGGCCCTCGGCCAGGA
CACCGCCCCAGGTCGTGCAGCCCAGCCCCAGCCTCAGCACGTTCTTCGGAGACACGGCCGCCAGCCACTCCTTGG
CCTCGGACTTCTTCGACTCCTTTACTACCTCCGCCTTCATTCCGTCAGCAATCCCGGCGCGGGCTCCCCGGCCC
CCGCCAGCCCCGCTCCCCCTCGCTGTGCCGGGACCGAGGGGCGCCCCGAACCCGTGGCCATGCGAGGGCCCCAGG
CAGTTCGCCCCCGCGTCGCCAGAGCCTTTCGCGCACATCCAGGCAGTGTTCGAGGGAGTGACGACCCCTTTG
CCACCGCCCTGAGCATGAGCGAGATGGACCGAGGAACGACGCCTGGCTTCCCGGCGAGGCTACGCGTGGAGTCC
TGCGGGCCGTGGCCACCCAGCAGCGCGGCGCGGTGTTCTGTTGACAAGGAGAACCTCACCATGCCGGGCCCTCAGGT
TCGACAACATCCAGGGAGATGCAGTTAAAGACTTGATCTTCGCTTTCGGGTGAAAAAGCTGCAGCAAAGAGAC
AAGTCTAAATGCCGACTCAGTGAACAATCTTTTGTGGATTGAAACAGCTAATCAGCTGCAGAACTGGAGGG
CAGCAGTGGACCTGTGCGGACGTCTCCTCACAGCCCACGGCCAGGGCTACGGCAAGAGCGGGCTGCTCACCAGCC
ACACGACAGATTCACTGCAGCTCTGGTTTGTGAGGCTGGCACTACTAGTGAAGTTGGGCCTTTTCCAGAATGCTG
AGATGGAATTTGAACCTTCGGAAATCTTGATCAGCCAGATCTTTATTACGAGTACTACCCGCACGTGTACCCTG
GGCGCAGGGGCTCCATGGTCCCCCTTCTCGATGCGCATCTTGACGCGGAGCTCAGCAGTACCTGGGGAACCCAC
AGGAGTCGCTGGATAGACTGCACAAGGTGAAGACTGTCTGCAGCAAGATCCTGGCCAATTTGGAGCAAGGCTTAG
CAGAAGACGGCGGCATGAGCAGCGTGACTCAGGAGGGCAGACAAGCCTCTATCCGGCTGTGGAGGTCACGTCTGG
GCCGGGTGATGTACTCCATGGCAAACGTGTCTGCTCCTGATGAAGGATTATGTGCTGGCCGTGGATGCGTATCATT
CGGTTATCAAGTATTACCCAGAGCAAGAGCCCCAGCTGCTCAGCGGCATCGGCCGATTTCCTCCAGATTGGAG
ACATAAAAAACAGCTGAAAAGTATTTTCAAGACGTTGAGAAAGTAACACAGAAATTAGACGGACTACAGGGTAAAA
TCATGGTTTTTGATGAACAGCGCGTTCCCTCACCTCGGGCAGAATAACTTTGCAGAAGCCACAGGTTCTTCACAG
AGATCTTAAGGATGGATCCAAGAAACGCAGTGGCCAACAACAACGCTGCCGTGTGTCTGCTCTACCTGGGCAAGC
TCAAGGACTCCCTGCGGCAGCTGGAGGCCATGGTCCAGCAGGACCCAGGCACTACCTGCACGAGAGCGTGCTCT
TCAACCTGACCACCATGTACGAGCTGGAGTCTCACGGAGCATGCAGAAGAAACAGGCCCTGCTGGAGGCTGTCTG
CCGCAAGGAGGGGGACAGCTTCAACACACAGTGCCTCAAGCTGGCCTAGCTGCCTCCAACACACTACGTACAGAA
GGACCCGGGTCTTTGAAACTGTGTCTTGAAGCTAATGTATTAATGTGACATGGAGGAACTCAATAAACTCCTGC
TTCAAAAAAAAAAAAAAAAAAAAAA

695/6881

FIGURE 648

MLRFLGEKAAAKRQVLNADSVEQSFVGLKQLISCRNWRAAVDLCGRLLTAHGQGYGKSGLLTSHTTDSLQLWFVR
LALLVKLGLFQNAEMEFEPFGNLDQPDLYEYYPHVYPGRRGSMVPFSMRILHAELQQYLGNPQESLDRHLHKVKT
VCSKILANLEQGLAEDGGMSSVTQEGRQASIRLWRSRLGRVMYSMANCLLLMKDYVLAVDAYHSVIKYYPEQEPQ
LLSGIGRISLQIGDIKTAEKYFQDVEKVTQKLDGLQGKIMVLMNSAFLHLGQNNFAEAHRFFTEILRMDPRNAVA
NNNAAVCLLYLGKLDKSLRQLEAMVQQDPRHYLHESVLFNLTTMYELESSRSMQKKQALLEAVAGKEGDSFNTQC
LKLA

696/6881
FIGURE 649

AATTCTGGCACGAGAAGACTTCCAGTTTGGAGTCGTTTGCTGCGGGGAGGGAATGAATGGGCGCTGGGAACACGCC
CGCGAGGTGGGGACGCGCCGGCCGTAGCGAGGTCCTTAGCGTGTGAGTGGCCGGGGTCGGGTGCGTTCCCCGCAG
CATGGAGGACGATGCACCACTGATCTACGGGCTGGAGTTCCAGGCACGTGCCTTAACACCTCAAACCTGCAGAAAC
AGATGCCATTCGGTTTTTGGTTGGGACGCAGTCTCTTAAATATGATAATCAGATCCATATCATAGATTTTGACGA
TGAAAACAACATTATAAAATAAAATGTCTCTCCATCAAGCGGGTGAAATCTGGCATATTAGCGCTAGCCCTGC
AGACAGAGGTGTGCTGACGACCTGCTACAACAGAACTTCAGACAGCAAAGTCTGACATGTGCAGCCGTGTGGAG
GATGCCGAAGGAATTGGAATCAGGCAGCCACGAGTCCCCTGATGATTTCATCCAGCACTGCACAGACCCCTGGAGCT
GCTCTGTACCTTGACAACACAGCCCATGGCAACATGGCCTGTGTCGTGTGGGAGCCAATGGGAGATGGGAAGAA
AATCATTTTCCTTGGCTGATAACCATATCCTGCTGTGGGATTTACAGGAAAGCTCGAGCCAGGCTGTGCTGGCCAG
CTCAGCGTCCCTGGAAGGGAAGGGACAACCTGAAGTTCACCTCAGGACGGTGGAGCCCACATCATAACTGCACCCA
GGTGGCCACAGCGAACGACACCACTCCGTTGGCTGGGACACCCGGAGCATGAGCCAGATCTACTGCATAGAGAA
TGCCACGGACAGCTGGTGGGGACCTTGACTTTAATCCCAATAAGCAGTACTACTTGGCCAGCTGCGGAGACGA
CTGTAAGGTGAAGTTCTGGGACACCCGAAATGTCAACGAACCCGTGAAGACCCTGGAGGAGCACTCCCACTGGGT
GTGGAACGTCCGCTACAACCACTCTCATGACCAGCTGGTCCTCACGGGCAGCAGTGACAGCAGAGTCATCCTTTC
CAACATGGTGTCCATCTCGTCCGAGCCCTTCGGCCACTTGGTAGACGACGATGACATCAGTGACCAGGAGGACCA
CCGTTCTGAAGAGAAGAGCAAGGAGCCCTGCAGGACAACGTGATCGCCACCTACGAGGAGCACGAGGACAGCGT
CTATGCCGTGGACTGGTCCTCGGCTGACCCGTGGCTGTTTGCCCTCCCTGAGCTATGACGGGAGGCTCGTGATCAA
CAGGGTGCCAGGGCCCTGAAGTACCACATCCTGCTA**TGA**CTCCCGGGCCTGGGTTATCCAGGTCCCATTGAGTG
GTTTTCTCTTGGCAGATTCTCAAACAGTCGCAGCTCTTGGAGGTGACTCGTGTTCAGGTGGATCCCTCTCTG
GGAGAGCCGCTGTTCCCTTCCTGTAGCAGCAGCATTATGAATGGGGTGAATGGGGCTATTGTCGACGGCACAGC
TAATGCCCCGAACCCAGCCCTGTGCGCAGAGACAGAGCCCCACATTATTATGTGAATAACAATGTTTTCTGTTTT
AAGGGTGTGAGGATTTTCGTTTTTAAAAAATGTCTGTTCTGCAGTAGTAACCTCTTCTTCTCTGAGAGTAA
AAAATGAAATAAAATAAATCCACGCTGACAAAAA

697/6881

FIGURE 650

MEDDAPVIYGLEFQARALTPQTAETDAIRFLVGTQSLKYDNQIHIIDFDDENNIINKNVLLHQAGEIWHISASPA
DRGVLTTTCYNRTSDSKVLTCAAVWRMPKELESGSHESPDDSSSTAQTLELLCHLDNTAHGNMACVVWEPMGDGKK
IISLADNHILLWDLQESSSQAVLASSASLEGKGQLKFTSGRWSPHHNCTQVATANDTTLRGWDTRSMSQIYCIEN
AHGQLVRDLDFNPNKQYYLASCDDCKVKFWDTRNVTEPVKTL EEHSHVWVNVRYNHS HDQLVLTGSSDSRVILS
NMVSISSEPFGLVDDDDISDQEDHRSEEKSKEPLQDNVIATYEEHEDSVYAVDWSSADPWLFASLSYDGRLVIN
RVPRALKYHILL

698/6881
FIGURE 651

CCCAGGCGCAGCCAATGGGAAGGGTCGGAGGCATGGCACAGCCAATGGGAAGGGCCGGGGCACCAAAGCCAATGG
GAAGGGCCGGGAGCGCGCGGGAGATTTAAAGGCTGCTGGAGTGAGGGGTCGCCGTGCACCCTGTCCCAG
CCGTCTGTCTGGCTGCTCGCTTCTGCTGCGCTCCACTATGCTCTCCCTCCGTGTCCCCTGCGGCCA
TCACGGACCCGACGAGCTGCTGCGCGCTGAAGGGCTCAGCTTGGTCGACAAGGAGAACACGCCGCCGG
CCCTGAGCGGGACCCGCTGCTGCGCCAGCAAGACCGGAGGAGGATCTTCCAGGAGCCACGGAGCCGAAAACTA
AAGCAGCTGCCCCCGGCTGGAGGATGAGCCGCTGCTGAGAGAAAACCCCGCGCTTTGTCTCTTCCCCATCG
AGTACCATGATATCTGGCAGATGTATAAGAAGGCAGAGGCTTCCTTTTGGACCGCCGAGGAGGTTGACCTCTCCA
AGGACATTCAGCACTGGGAATCCCTGAAACCCGAGGAGAGATATTTATATCCCATGTTCTGGCTTTCTTTGCAG
CAAGCGATGGCATAGTAAATGAAACTTGGTGGAGCGATTTAGCCAAGAAGTTCAGATTACAGAAGCCCGCTGTT
TCTATGGCTTCCAAATTGCCATGGAAAACATACATTCTGAAATGTATAGTCTTCTTATTGACACTTACATAAAAG
ATCCCAAAGAAAGGGAATTTCTTCAATGCCATTGAAACGATGCCTTGTGTCAAGAAGAAGGCAGACTGGGCCT
TGCGCTGGATTGGGGACAAAGAGGCTACCTATGGTGAACGTGTTGTAGCCTTTGCTGCAGTGGAAGGCATTTCT
TTTCCGGTTCTTTTGCCTGATATTCTGGCTCAAGAAACGAGGACTGATGCCTGGCCTCACATTTTCTAATGAAC
TTATTAGCAGAGATGAGGGTTTACACTGTGATTTTGGCTTGCCTGATGTTCAAACACCTGGTACACAAACCATCGG
AGGAGAGAGTAAGAGAAATAATTATCAATGCTGTTCCGATAGAACAGGAGTTCCTCACTGAGGCCTTGCCTGTGA
AGCTCATTGGGATGAATTGCACTCTAATGAAGCAATACATTGAGTTTGTGGCAGACAGACTTATGCTGGAACCTGG
GTTTTAGCAAGGTTTTTCAGAGTAGAGAACCCATTTGACTTTATGGAGAATATTTCACTGGAAGGAAAGACTAACT
TCTTTGAGAAGAGAGTAGGCGAGTATCAGAGGATGGGAGTGATGTCAAGTCCACAGAGAATTCTTTTACCTTGG
ATGCTGACTTCTAAATGAAGTGAAGATGTGCCCTTACTTGGCTGATTTTTTTTTTCCATCTCATAAGAAAAATCA
GCTGAAGTGTTACCAACTAGCCACACCATGAATTGTCCGTAATGTTTATTAAACAGCATCTTTAAACTGTGTAGC
TACCTCACAACCAGTCTCTGTCTTTATAGTGCTGGTAGTATCACCTTTTGGCAGAAGGCCTGGCTGGCTGTGAC
TTACCATAGCAGTGACAATGGCAGTCTTGGCTTTAAAGTGAGGGGTGACCCTTTAGTGAGCTTAGCACAGCGGGA
TTAAACAGTCTTTTAAACAGCACAGCCAGTTAAAGATGCAGCCTCACTGCTTCAACGCAGATTTTAAATGTTTAC
TTAAATATAAACCTGGCACTTTACAAACAAATAAACATTGTTTGTACTCACGGCGGCGATAATAGCTTGATTTA
TTTGGTTTCTACACCAAATACATTCTCCTGACCATAATGGGAGCCAATTCACAATTCACTAAGTGACTAAAGTA
AGTTAACTTGTGTAGACTAAGCATGTAATTTTTAAGTTTTATTTTAAATGAATTAATAATTTTGTAAACCACTT
TAAAGTCAGTCTGTGTATACCTAGATATTAGTCAGTTGGTGCCAGATAGAAGACAGGTTGTGTTTTATCCTGT
GGCTTGTGTAGTGCTCTGGGATTCTCTGCCCCCTCTGAGTAGAGTGTTGTGGGATAAAGGAATCTCTCAGGGCAA
GGAGCTTCTTAAAGTTAAATCACTAGAAATTTAGGGGTGATCTGGGCCTTCATATGTGTGAGAAGCCGTTTCATT
TATTTCTCACTGTATTTTCTCAACGCTCTGGTTGATGAGAAAAAATCTTGAAGAGTTTTCATATGTGGGAGCTA
AGGTAGTATTGTAAATTTCAAGTCATCCTTAAACAAAATGATCCACCTAAGATCTTGCCCTGTAAAGTGGTGA
AATCAACTAGAGGTGGTTCTTACAAGTTGTTTATTCTAGTTTTGTTTGGTGTAAAGTAGGTTGTGTGAGTTAATC
ATTTATATTTACTATGTCTGTTAAATCAGAAATTTTTATTATCTATGTTCTTCTAGATTTTACCTGTAGTTTCA
AAAAAAAAAAAAAAAAAAAAAAAAA

699/6881
FIGURE 652

MLSLRVPLAPITDPQQQLSPLKGLSLVDKENTPPALSGTRVLASKTARRIFQEPTEPKTKAAAPGVEDEPLLRE
NPRRFVIFPIEYHDIWQMYKKAEASFWTAEVDLSKDIQHWESLKPEERYFISHVLAFFAASDGIVNENLVERFS
QEVQITEARCFYGFQIAMENIHSEMYSLIDITYIKDPKEREFLFNAIETMPCVKKKADWALRWIGDKEATYGERV
VAFAAVEGIFFSGSFASIFWLKKRGLMPGLTFSNELISRDEGLHCDFACLMFKHLVHKPSEERVREIIINAVRIE
QEFLTEALPVKLIGMNCTLMKQYIEFVADRLMLELGF SKVFRVENPFD FMENISLEGKTNFFEKRVG EYQRMGVM
SSPTENSFTLDADF

700/6881
FIGURE 653

GTCGCTGAGGCGCCCCATGGCCTTCGCCCCGCGGCTCCTGCGCGGGCCACTGTCGGGGCCGCTGCTCGGGCGGGCGC
GGGGTCTGCGCTGGGGCCATGGCTCCGCCGCGCCGCTTCGTCTGGAGCTTCCCGACTGCACCCTGGCTCACTTC
GCCCTAGGCGCCGACGCCCCGCGCAGCAGACGCCCCGACCCCCGCTGGCGGGCGCTGCTGGGGCCCCCGGAG
CGCAGCTACTCGCTGTGCGTGCCCCGTGACCCCGACGCGGCTGCGGGGCCGGGTCCGGGCGGCGCGGCTGCAC
CAGCGCCTGCTGCACCAGCTGCGCCGCGGCCCCCTTCCAGCGGTGCCAGCTGCTCAGGCTGCTCTGCTACTGCCCG
GGCGGCCAGGCCGCGCGCACAGCAAGGCTTCCTGCTGCGCGACCCCCCTGGATGACCCTGACACCCGGAAGCG
CTGCTCGAGCTGCTGGGCGCCTGTGAGGAGGACACGCCCCGCACTTGGGCGAGTTCGAGGCCGACCCGCGCGGC
CAGCTGTGGCAGCGCCTCTGGGAGGTGCAAGACGGCAGGCGGCTGCAGGTGGGCTGCGCACAGGTCTGCCCCGT
CCGGAGCCCCCGCTGCACCCGGTGGTGCCAGACTTGCCCCAGTTCCGTGGTCTTCCCGACCGGAAGCCGCCCGG
GCCGTTTTGGAGGAGTGTACCTCCTTTATTCCCTGAAGCCCGGCAGTGCTTGACCTGGTCGACCACTGCCAAAA
CAGATCCAGAAAGGAAAGTTCCAGGTTGTTGCCATCGAAGGACTGGATGCCACGGGTAAAACCAGGTGACCCAG
TCAGTGGCAGATTCACTTAAGGCTGTCTCTTAAAGTCACCACTCTTGCATTGGCCAGTGGAGGAAGATCTTT
GATGATGAACCAACTATCATTAGAAGAGCTTTTACTCTTTGGGCAATTATATTGTGGCCTCCGAAATAGCTAAA
GAATCTGCCAAATCTCCTGTGATTGTAGACAGGTACTGGCACAGCACGGCCACCTATGCCATAGCCACTGAGGTG
AGTGGGGGTCTCCAGCACCTGCCCCAGCCCATCACCTGTGTACCACTGGCCAGAGGACCTGCTCAAACCTGAC
CTTATCCTGCTGCTCACTGTGAGTCTGAGGAGAGGTGTCAGAGGCTGCAGGGCCGGGCGATGGAGAAGACCAGG
GAAGAAGCAGAACTTGAGGCCAACAGTGTGTTTCGTCAAAAGGTAGAAATGTCTACCAGCGGATGGAGAATCCT
GGCTGCCATGTGGTTGATGCCAGCCCCCTCCAGAGAAAAGTCTGCAGACGGTATTAAAGCCTAATCCAGAATAGT
TTTAGTGAACCGTAGTTACTCTGGCCAGGTGCCACGTCTAAGTAGATTAGATGTTGTTTGAACATCTACATCCA
CCATTTGTTATGCAGTGTTCCTCAAAATTTCTGTTCTACAAGCATGTTGTGTGGCAGAAAAGTGGAGACAGGCATC
TTAATTTTACTTCAGCCATCGTACCCTCTTCTGACTGATGGACCCGTCATCACAAAGTCCCTCTCATCATGTTT
CAGTGAGAGGCCAGCGATTGCTTTCTTCTGGCATAGTAAACATTTCTTGAACATATGTTTCACTTAATCACT
ACCAATATCTGGAAGACCTGTCTTACTCAGACAGCACAGGTGTACAGAAGCAGCAGACAAGATCTTCCAGATC
AGCAGGGAGACCCCGGAGCCTCTGCTTCTCCTACTGGCATGTGATGAGATCGTGACATGCCACATTGGCTT
CTTCCACATCTGGTTGCACTCGTCATGATGGGCTCGCTGCATCTCCCTCAGTCCCAATTCAGAGCCAAGTGT
CCTGCAGAGGCTGTCTATGTGCTGAGGCTCAGAGCCCTTTGATAGGCTTCTGAGTCATATATAAAGACATTC
CAAAGAAGGCAATTGATCTTGTGCTGAGGCTCAGAGCCCTTTGATAGGCTTCTGAGTCATATATAAAGACATTC
AAGCCAAGATGCTCCAAGTGAATATAACCAACCTTCTCTGAATTATATTTGCTTATTTATTTCTTTCTTT
TTTTCTAAAGTATGGCTCTGAATAGAATGCACATTTTCCATTGAACTGGATGCATTTCAATTAGCCAATCCAGTA
ATTTATTTATATTAATCTATACATAATATGTTTCCCTCAGCATAGGAGCTATGATTCAATTAATTAAGTGGAGTC
AAAACGCTAAATGCAATGTTTGTGTTGTTGTTTTCATTACACAACTTAATTTGTCTTGTAAATAAGTACAGTGG
ATCTTTGAGTGGGATTTCTTGGTAAATATCTTGCACCTGAATGTCTCATGATTACATATGAAATCGCTTTGACA
TATCTTTAGACAGAAAAAGTAGCTGAGTGAGGGGAAATATAGAGCTGTGTGACTTTAGGGAGTAGGTTGAAC
CAGGTGATTACCTAAAATTCCTTCCAGTTCAAAGGCAGATAAATCTGTAAATTAATTTATCTATCTACCATTTT
TTAAGAAGACATTACTCCAAAATAATTAATTTAAGGCTTATCAGGTCTGCATATAGAATCTTAAATCTAATA
AAGTTTCATGTTAATGTCATAGGATTTTAAAAGAGCTATAGGTAATTTCTATATAATATGTGTATATTAAATG
TAATTGATTTAGTTGAAAGTATTTTAAAGCTGATAAATAGCAATTAGGGTTCTTTGCAATGTGGTATCTAGCTGT
ATTATTGGTTTTATTACTTTAAACATTTTGAAGCTTACTGGCAGCCTAGAAAAACAAACAATTAATGTAT
CTTTATGTCCCTGGCACATGAATAAACTTTGCTGTGGTTTACT

701/6881

FIGURE 654

MAFARRLLRGPLSGPILLGRRGVCAGAMAPRRFVLELPDCTLAHFALGADAPGDADAPDPRLAALLGPPERSYSL
CVPVTPDAGCGARVRAARLHQRLHQLRRGPFQRCQLRLLCYCPGGQAGGAQQGFLLRDPLDDPDTRQALLELL
GACQEAPRPHLGEFEADPRGQLWQRLWEVQDGRRLQVGCAQVVPVPEPPLHPVVPDLPSSVVFPDREAARAVLEE
CTSFIEPEARAVLDLVDQCPKQIQKGKFQVVAIEGLDATGKTTVTQSVADSLKAVLLKSPPSCIGQWRKIFDDEPT
IIRRAFYSLGNYIVASEIAKESAKSPVIVDRYWHSTATYAIATEVSGGLQHLPPAHHVPYQWPEDLLKPDILILL
TVSPEERLQRLQGRGMEKTREEAELEANSVFRQKVMSYQRMENPGCHVVDASPSREKVLQTVLSLIQNSFSEP

702/6881
FIGURE 655

GTGGTGGGACTCGCGTCGCGGCCGCGGAGACGTGAAGCTCTCGAGGCTCCTCCCGCTGCGGGTCGGCGCTCGCCC
TCGCTCTCCTCGCCCTCCGCCCCGGCCCCGGCCCCGCGCCCGCCATGGAGAAGACTGAGCTGATCCAGAAGGCCA
AGCTGGCCGAGCAGGCCGAGCGCTACGACGACATGGCCACCTGCATGAAGGCAGTGACCGAGCAGGGCGCCGAGC
TGTCCAACGAGGAGCGCAACCTGCTCTCCGTGGCCTACAAGAACGTGGTCGGGGGCCGAGGTCGCGCTGGAGGG
TCATCTCTAGCATCGAGCAGAAGACCGACACCTCCGACAAGAAGTTGCAGCTGATTAAGGACTATCGGGAGAAAG
TGGAGTCCGAGCTGAGATCCATCTGCACCACGGTGTGGAATTGTTGGATAAATATTTAATAGCCAATGCAACTA
ATCCAGAGAGTAAGGTCTTCTATCTGAAAATGAAGGGTGATTACTTCCGGTACCTTGCTGAAGTTGCGTGTGGTG
ATGATCGAAAACAAACGATAGATAATTCCCAAGGAGCTTACCAAGAGGCATTGATATAAGCAAGAAAGAGATGC
AACCACACACCCAATCCGCTGGGGCTTGCTCTTAACCTTTTCTGTATTTTACTATGAGATTCTTAATAACCCAG
AGCTTGCCTGACGCTGGCTAAAACGGCTTTTGATGAGGCCATTGCTGAACTTGATACACTGAATGAAGACTCAT
ACAAAGACAGCACCTCATCATGCAGTTGCTTAGAGACAACCTAACACTTTGGACATCAGACAGTGCAGGAGAAG
AATGTGATGCGGCAGAAGGGGCTGAAAACTAAATTCCATACAGGGTGTCACTCCTTCTTTCCTTCAAGAAACCTTTT
TACACATCTCCATTCTTATTCACCTTGGATTTCTATAGCAAAGAAACCCATTTCATGTGTATGGAATCAACTGT
TTATAGTCTTTTACACTGCAGCTTTGGGAAAACCTTCTTCTTGTATTGTGTTTGTCTTGGCCTTCTCGGTGTG
CAGTACTGCTGTAGAAAAGTATTAATAGCTTCATTTCATATAAACATAAGTAACCTCCCAAACACTTATGTAGAGG
ACTAAAAATGTATCTGGTATTTAAGTAATCTGAACCGATTCTGCAAGTGAAGTGTGTTTTGTATTACTGTGAAAAT
AAGAAAATGTAGTTAATTACAATTTAAAGAGTATTCACATAACTTCTTAATTTCTACATTCCCTCCCTTACTCT
TCGGGGGTTTCTTTTCAGTAAGCAACTTTTCCATGCTCTTAATGTATTCTTTTGTAGTAGGAATCCGGAAGTATT
AGATTGAATGGAAAAGCACTTGCCATCTCTGTCTAGGGGTCACAAATTGAAATGGCTCCTGTATCACATACGGAG
GTCTTGTGTATCTGTGGCAACAGGGAGTTTCTTATTCACTCTTTATTTGCTGCTGTTTAAGTTGCCAACCTCCC
CTCCCAATAAAAAATTCACCTTACACCTCCTGCCTTTGTAGTTCTGGTATTCACTTTACTATGTGATAGAAGTAGCA
TGTTGCTGCCAGAATACAAGCATTGCTTTTGGCAAATTAAAGTGCATGTCATTTCTTAATACACTAGAAAGGGGA
AATAAATTAAAGTACACAAGTCCAAGTCTAAACCTTTAGTACTTTTCCATGCAGATTGTGCACATGTGAGAGGG
TGTCAGTTTGTCTAGTGATTGTTATTTAGAGAGTTGGACCACTATTGTGTGTTGCTAATCATTGACTGTAGTCC
CAAAAAAGCCTTTGTGAAAATGTTATGCCCTATGTAACAGCAGAGTAACATAAAATAAAAGTACATTTTATAAACC
ATTTACTATGGCTTTGTAACAATTGCATACCCATATTTTAAAGGGACAGGTGAATTTACTACTTTCTAAAGTTTAT
TGATACTTCCCTTTTATGTAAAATGTAGTAGTGATACCTATATTTCCACATTGTGCATTGTGACACACTTGTCTA
GGGATGCCTGGAAGTGATAAAATTGGACTGCATTTCTTAGAGTGTTTTACTATAGATCAGTCTCATGGGCCATC
TCTTCTCAGATGTAAATGATATCTGGTTAAGTGTTATATGGAATAAAGTGGACATTTTAAACCTA

703/6881
FIGURE 656

MEKTELIQKAKLAEQAERYDDMATCMKAVTEQGAELSNEERNLLSVAYKNVVGRRSAWRVISSIEQKTDTSDDK
LQLIKDYREKVESELRSICTTVLELLDKYLIANATNPESKVFYLMKGDYFRYLAEVACGDDRQQTIDNSQGAYQ
EAFDISKKEMQPTHPIRLGLALNFSVFYIEILNNPELACTLAKTAFDEAIAELDTLNEDSYKDSLIMQLLRDNL
TLWTSDSAGEECDAAEGAEN

704/6881
FIGURE 657

GGCACGAGGCTCCGGTGTGTCTGTCGGTTGCAGTGTGGAGGTCGGCGCCGGCCCCCGCCTTCCGCGCCCCCAC
GGGAAGGAAGCACCCCCGGTATTAAACGAACGGGGCGGAAAGAAGCCCTCAGTCGCCGGCCGGGAGGCGAGCCG
ATGCCGAGCTGTCCACGTCCACCATGCCGGGCATGATCTGCAAGAAGCCAGACCTCGAGTTTGACTCGCTACAG
CCCTGCTTCTACCCGGACGAAGATGACTTCTACTTCGGCGGCCCGACTCGACCCCCCGGGGAGGACATCTGG
AAGAAGTTTGAGCTGCTGCCCACGCCCCCGCTGTGCCCCAGCCGTGGCTTCGCGGAGCACAGCTCCGAGCCCCCG
AGCTGGGTACGGAGATGCTGCTTGAGAACGAGCTGTGGGGCAGCCCGGCCGAGGAGGACGCGTTCGGCCTGGGG
GGACTGGGTGGCTCACCCCCAACCCGGTCATCTCCAGGACTGCATGTGGAGCGGCTTCTCCGCCCGCGAGAAG
CTGGAGCGCGCCGTGAGCGAGAAGCTGCAGCACGGCCGCGGGCCGCCAACCGCCGGTTCACCGCCCCAGTCCCCG
GGAGCCGGCGCCGACGCCCTGCGGGTTCGCGGGCAGCGCGGGGCTGCGGGAGCCGGCCGCGCGGGGCCGCCCTG
CCCGCCGAGCTCGCCCCACCCGGCCGCGGAGTGCGTGGATCCCGCCGTGGTCTTCCCCTTTCCCGTGAACAAGCGC
GAGCCAGCGCCCGTGCCCGCAGCCCCGCGCAGTGCCCCGGCGGGCCCTGCGGTGCGCTCGGGGGCGGGTATT
GCCGCCCAGCCGGGGCCCCGGGGTTCGCCCTCCGCGCCAGGCGGCCGCCAGACCAGCGGGCGACCACAAG
GCCCCAGTACCTCCGGAGAGGACACCTGAGCGATTGAGATGATGAAGATGATGAAGAGGAAGATGAAGAGAA
GAAATCGACGTGGTCACTGTGGAGAAGCGGCGTTCCTCCTCCAACACCAAGGCTGTCAACACATTCACCATCACT
GTGCGTCCCAAGAACGAGCCCTGGGTCCCGGGAGGGCTCAGTCCAGCGAGCTGATCCTCAACGATGCCTTCCC
ATCCACGAGCAGCACAACTATGCCGCCCTCTCCTACGTGGAGAGTGAGGATGCACCCACAGAGAAGAAGATA
AAGAGCGAGGCGTCCCCACGTCCGCTCAAGAGTGTATCCCCCAAAGGCTAAGAGCTTGAGCCCCCGAACTCT
GACTCGGAGGACAGTGAGCGTCGCAGAAACCACAACATCCTGGAGCGCCAGCGCCGCAACGACCTTCGGTCCAGC
TTTCTACGCTCAGGGACCAGTGCCGGAGTTGGTAAAGAATGAGAAGGCCGCCAAGGTGGTCATTTTGAAAAAG
GCCACTGAGTATGTCCACTCCCTCCAGGCCGAGGAGCACCAGCTTTTGCTGGAAAAGGAAAAATTGCAGGCAAGA
CAGCAGCAGTTGCTAAAGAAAATTGAACACGCTCGGACTTGCTAGACGCTTCTCAAAACTGGACAGTCACTGCCA
CTTTGCACATTTTGATTTTTTTTTTAAACAAACATTGTGTGACATTAGAATGTTGGTTTACTTTCAAATCGGT
CCCCTGTCGAGTTCCGGCTCTGGGTGGGCAGTAGGACCACCACTGTGGGGTTCGTGCTGGGACCTTGAGAGCCTGC
ATCCCAGGATGCTGGGTGGCCCTGCAGCCTCCTCCACCTCACCTCCATGACAGCGCTAAACGTTGGTGACGGTTG
GGAGCCTCTGGGGCTGTTGAAGTCACCTTGTTGTTTCCAAGTTTCCAACAACAGAAAGTCATTCCCTTCTTTTTA
AAATGGTGCTTAAGTTCCAGCAGATGCCACATAAGGGGTTTGCCATTTGATACCCCTGGGGAACATTTCTGTAAA
TACCATTGACACATCCGCCTTTTGATACATCCTGGGTAATGAGAGGTGGCTTTTGCGGCCAGTATTAGACTGGA
AGTTTACCTAAGTACTGTAATAATACCTCAATGTTTGAGGAGCATGTTTTGTATACAAATATATTGTTAATCT
CTGTTATGTACTGTACTAATTCTTACACTGCCGTGATACCTTAGTATGACGCTGATACATAACTAAATTTGATAC
TTATATTTTCGTATGAAAATGAGTTGTGAAAGTTTTGAGTAGATATTACTTTATCACTTTTTGAACTAAGAACT
TTTGTAAGAAATTTACTATATATATATGCTTTTTCCTAGCCTGTTTCTTCTGTTAATGTATTGTTCATGTT
TGGTGCAAGAACTGGGTAAATGCAAGTTCTGTGTTTAAATTTCTTCAAATGTATATATTTAGTGTGCATCTT
ATAGCACTTTGAAATACCTCATGTTTATGAAAATAAATAGCTTAAATTAAAAAAAAAAAAAAAAAA

705/6881
FIGURE 658

MPSCSTSTMPGMICKNPDLEFDSLQPCFYDDEDDFYFGGPDSTPPGEDIWKKFELLPTPPLSPSRGFAEHSSEPP
SWVTEMLLENELWGSPAEDAFGLGGLGGLTPNPVILQDCMWSGFSAREKLERAVSEKLQHGRGPPTAGSTAQSP
GAGAASPAGRGHGGAAGAGRAGAALPAELAHPAECVDPVAVFFFPVNKREPAPVPAAPASAPAAGPAVASGAGI
AAPAGAPGVAPPRPGGRQTSGGDHKALSTSGEDTSDSDDEDDEEEDDEEEDIDVVTVEKRRSSSNTKAVTTFTIT
VRPKNAALGPGRAQSSELILKRCLPIHQHNYAAPSPYVESEDAPPQKKIKSEASPRPLKSVIPPKAKSLSPRNS
DSEDSERRRNHNILERQRRNDLRSSFLTLDHVPPELVKNEKAAKVVLKKATEYVHSLQAEHQLLLEKEKLQAR
QQLLKKIEHARTC

706/6881
FIGURE 659

GTCAGTCCCTCCTGTAGCCGCCGCCGCCGCCGCCGCCCTCTGCCAGCAGCTCCGGCGCCACCTCGGGCCG
GCGTCTCCGGCGGGCGGGAGCCAGGCGCTGACGGGCGCGGGCGGGCGGGCGAGCGCTCCTGCGGCTGCGACTCA
GGCTCCGGCGTCTGCGCTTCCCCATGGGGCTGGCCTGCGGCGCCTGGGCGCTCTGAGATTGTCACTGCTGTTCCA
AGGGCACACGCAGAGGGATTGGAATTCCTGGAGAGTTGCCTTTGTGAGAAGCTGGAAATATTTCTTTCAATTCC
ATCTCTTAGTTTTCCATAGGAACATCAAGAAATCATGAACAACCTTTGGTAATGAAGAGTTTGACTGCCACTTCCT
CGATGAAGGTTTTACTGCCAAGGACATTCTGGACCAGAAAATTAATGAAGTTTCTTCTCTGATGATAAGGATGC
CTTCTATGTGGCAGACCTGGGAGACATTCTAAAGAAACATCTGAGGTGGTTAAAAGCTCTCCCTCGTGTCACCCC
CTTTTATGCAGTCAAATGTAATGATAGCAAAGCCATCGTGAAGACCCTTGCTGCTACCGGGACAGGATTTGACTG
TGCTAGCAAGACTGAAATACAGTTGGTGCAGAGTCTGGGGGTGCCTCCAGAGAGGATTATCTATGCAAATCCTTG
TAAACAAGTATCTCAAATTAAGTATGCTGCTAATAATGGAGTCCAGATGATGACTTTTGATAGTGAAGTTGAGTT
GATGAAAGTTGCCAGAGCACATCCCAAAGCAAAGTTGGTTTTGCGGATTGCCACTGATGATTTCCAAAGCAGTCTG
TCGTCTCAGTGTGAAATTCGGTGCCACGCTCAGAACCAGCAGGCTCCTTTTGGAACGGGCGAAAGAGCTAAATAT
CGATGTTGTTGGTGTGTCAGCTTCCATGTAGGAAGCGGCTGTACCGATCCTGAGACCTTCGTGCAGGCAATCTCTGA
TGCCCGCTGTGTTTTGACATGGGGGCTGAGGTTGGTTTCAGCATGTATCTGCTTGATATTGGCGGTGGCTTTCC
TGATCTGAGGATGTGAAACTTAAATTTGAAGAGATCACCGCGTAATCAACCCAGCGTTGGACAAATACTTTCC
GTCAGACTCTGGAGTGAGAATCATAGCTGAGCCCCGCGAGATACTATGTTGCATCAGCTTTCACGCTTGCAAGTTAA
TATCATTGCCAAGAAAATTGTATTAAAGGAACAGACGGGCTCTGATGACGAAGATGAGTCGAGTGAGCAGACCTT
TATGTATTATGTGAATGATGGCGTCTATGGATCATTTAATTGCATACTCTATGACCACGCACATGTAAAGCCCCT
TCTGCAAAAGAGACCTAAACCAGATGAGAAGTATTATTCATCCAGCATATGGGGACCAACATGTGATGGCCTCGA
TCGGATTGTTGAGCGCTGTGACCTGCCTGAAATGCATGTGGGTGATTGGATGCTCTTTGAAAACATGGGCGCTTA
CACTGTTGCTGCTGCCCTCTACGTTCAATGGCTTCCAGAGGCCGACGATCTACTATGTGATGTCAGGGCCTGCGTG
GCAACTCATGCAGCAATTCAGAACCCCGACTTCCCACCGAAGTAGAGGAACAGGATGCCAGCACCCCTGCCTGT
GTCTTGTGCTGGGAGAGTGGGATGAAACGCCACAGAGCACCTGTGCTTCGGCTAGTATTAATGTGTAGATAGC
ACTCTGGTAGCTGTAACTGCAAGTTTAGCTTGAATTAAGGGATTGGGGGGACCATGTAACCTAATTACTGCTA
GTTTTGAAATGTCTTTGTAAGAGTAGGGTCGCCATGATGCAGCCATATGGAAGACTAGGATATGGGTCACACTTA
TCTGTGTTCCATGGAACATATTTGAATATTTGTTTTATATGGATTTTTTATCACTCTTCAGACACGCTACTCAA
GAGTGCCCCCTCAGCTGCTGAACAAGCATTGTAGCTTGTACAATGGCAGAATGGGCCAAAAGCTTAGTGTTGTGA
CCTGTTTTTAAATAAAGTATCTTGAATAATTAGGC

707/6881
FIGURE 660

MNFGNEEFDCHFLDEGFTAKDILDQKINEVSSSDDKDAFYVADLGDILKKHLRWL KALPRVTPFYAVKCND SKA
IVKTLAATGTGFDCASKTEIQLVQSLGVPPERIIYANPCKQVSQIKYAANNGVQMMTFDSEVELMKVARAHPKAK
LVLRIATDDSKAVCRLSVKFGATLRTSRLLLERAKELNIDVVGVSFHVSGSGCTDPETFVQAISDARCVFDMGAEV
GFSMYLLDIGGGFPGSEDVKLKFEETGVINPALDKYFPSDSGVRIIAEPGRYYVASAFTLAVNIIAKKIVLKEQ
TGSDDEDESSEQTFMYVNDGVYGSFNCILYDHAHVKPLIQKRPKPDEKYYSSSIWGPTCDGLDRIVERCDLPEM
HVGDWMLFENMGAYTVAAASTFNGFQRPTIYYVMSPAWQLMQQFQNPDPPEVEEQDASTLPVSCAWESGMKRH
RAACASASINV

708/6881
FIGURE 661

ATGAGTCAGCAGCGGCCGCGGAGGAAGTTACCCAGTCTCCTCCTGGACCCGACGGAGGAGACGGTTCGCCGTCGG
TGCCGAGACCCCATCAACGTGGAGGGCCTGCTGCCATCAAAAATAAGGATTAATTTAGAAGATAATGTACAATAT
GTGTCCATGAGAAAAGCTCTAAAAGTGAAGAGACCTCGTTTTGATGTATCGCTGGTTTATTTAACTCGAAAATTT
ATGGATCTTGTGAGATCTGCTCCCGGGGGTATTCTTGACTTAAACAAGGTTGCAACGAAACTGGGAGTCCGAAAG
CGGAGAGTGTATGACATCACCAATGTCTTAGATGGAATCGACCTCGTTGAAAAGAAATCCAAGAACCATATTAGA
TGGATAGGATCTGATCTTAGCAATTTTGGAGCAGTTCCCCAACAAAAGAAGCTACAGGAGGAACTTTCTGACTTA
TCAGCAATGGAAGATGCTTTGGATGAGTTAATTAAGGATTGTGCTCAGCAGCTGTTGAGTTAACAGATGACAAA
GAAAATGAAAGACTAGCATATGTGACCTATCAAGACATTCATAGCATTAGGCCCTCCATGAACAGATCGTCATT
GCAGTTAAAGCTCCAGCAGAAACCAGATTGGATGTTCCAGCTCCAGAGAAGACTCTATCACAGTGCACATAAGG
AGCACCAACGGACCTATCGATGTCTATTTGTGTGAAGTGGAGCAGGGTCAGACCAGTAACAAAAGGTCTGAAGGT
GTGGGACCTCTTCATCTGAGAGCACTCATCCAGAAGGCCCTGAGGAAGAAGAAAATCCTCAGCAAAGTGAAGAA
TTGCTTGAAGTAAGCAACTGATGGCATTGAGAATTTATGTATCACTGAGTTTTTGGGAATATCTTCGTGGAGA
ATTACGCATCAAATTTGATTCTCAGAGCAATAAATTATCCATGAAGTGCTCTCGTTCTCAGTAGCGGCATCATGG
CCAGTAGTGTCTTTGAGGAGTTCACCACCTAGATTACTGAGTAATTGTGGTTCCACATTTGAAAACAACCTCCTT
TTATAATTATTCAGTGTCTTTTGTGAGTGAATAGACATCTTGCCCTCCTGAAGTAGCTTCATCACAGAGTGTGAT
GAAGACAGACAGTCAGGCTGAAATGGACAGTTCTTTGTGGACTCTACCCTTCCCTTCAAGGAGTATGTCATATAT
CACAAAAGAAATTGCCCTTACACTGGTTCATGTTTGCAGTTACTGTTGTACATTGCATAGATGTACACACGAATTT
AAATGTGATGTCTTTGTATATATCTGTATAATGTTGAGATTACTTACGAAATATGTCTGAGTGACACTTTTCACC
CTTGTACAGCCAAAATAATGTATATATGGAAAGTGACAGACAAATTCTCTAATCTCTTTGGTACCTATAACTTAT
TAGAATCCTCTGGATGAGGGTTAGAAGAGACTTTTTCCAACTTCTACATGTAGAAGTATCATAAATGTGCTACA
CATTTATGTTTGTGGATTAAATTAAAGTATTTAATATGGTTTTTCAGTGCTAAAATTGGAGTCAGATACTTCTTG
GTTTTAAGCTGTCTACCTAATTGCTGTCTCCAGCAGACTGGTGGCATGCCAGTGGCTTTGGGGGCAAGGATAG
AAATGCCATCAGGAAATAGCTGAATTCATTGTGAACATGAATTCAGTCATGGTGATAATTGGAAACTCCTTTCA
GGTTTTTGCAAGTAGATTTGTAAATGTTTGTGTATGCAGCCTTGCTGTTGAGTCAGTCCAAGGGGTTTTACTTAG
GACAAGTTGTACCTTGCCCTCTCTCCAGCTGTGCTCCACATTTTCACATACCTAGCTGTTTCTACCTCATTGGG
TAAGTCATTTACCACCTGTGTCCTCAGTTTACTCTGTAGTTTACCATTAGACTGTGAGCTCCTTGAGGGACTTTG
TCATAATCACTGTTACATCCCAGTGCCTCACACCATGCCTGGCCCTTAAGAAGTGCTCAATAAATGTCTGAACAA
ATAA

709/6881
FIGURE 662

CTCTGTAGCTGTGACCCTGATACCGCGTGGTGTGCTCCGAACACATGGTGCCCGAAGCAAGGCGGCGTCCAGAA
GCCCTAGGTCCCAGAGGTCCGCTCAGCGGCAGGCGCATAAGGCGGGGCCGGCGCGGGCCTTTTCCTTCCATCGGAA
CCGTTCTCCCGGGGCTGAGTCCCTGCCCCGACTCCGAACGCCGAAGACCAGGGGCCGGAAGCGCGCGCCGCACT
GCCACGCCGTGTGTCAGTCGGGAGGGAGGCGAGCAGGCGAAGCCGCGGAGGACGGGGTGAAGATGGCGGCCCTTC
TCCGAGATGGGTGTAATGCCTGAGATTGCACAAGCTGTGGAAGAGATGGATTGGCTCCTCCCACTGATATCCAG
GCTGAATCTATCCCATTTGATCTTAGGAGGAGGTGATGTACTTATGGCTGCAGAAACAGGAAGTGGCAAACTGGT
GCTTTTAGTATTCCAGTTATCCAGATAGTTTATGAAACTCTGAAAGACCAACAGGAAGCAAAAAAGGAAAAACA
ACAATTAAACTGGTGCTTCAGTGCTGAACAAATGGCAGATGAACCATATGACAGAGGATCTGCTTTTGAATT
GGGTCAGATGGTCTTTGTTGTCAAAGCAGAGAAGTAAAGGAATGGCATGGGTGTAGAGCTACTAAAGGATTAAATG
AAAGGGAACACTACTATGAAGTATCCTGTCTATGACCAAGGGTTATGCAGGGTCGGGTGGTCTACCATGCAGGCC
TCTTTGGACCTAGGTACTGACAAGTTTGGATTGGCTTTGGTGGAAACAGGAAGAAATCCCATAAACAACTTT
GATAATTATGGAGAGGAATTCATATGCATGATACCATTTGGATGTTACCTGGATATAGATAAGGGACATGTCAAG
TTCTCCAAAAATGGAAAAGATCTTGGTCTGGCATTGAAATACCACCATATGAAAAACCAAGCCCTCTTTCTCT
GCCTGTGTTTTGAAGAATGCTGAAGTAAATTTAACTTCGGTGAAGAGGAATTTAAGTTTCCACCAAAAGATGGC
TTTGTGCTCTTTCCAAAGGCACCGGATGGTTACATTGTCAAATCACAGCACTCAGGTAATGCACAGGTGACACAA
ACAAAGTTTCTCCCAATGCTCCGAAAGCTCTCATTGTGAACCTTCCCGGGAGTTAGCTGAACAACTTTGAAC
AACATCAAGCAGTTTAAGAAATACATTGATAATCCTAAATTAAGGGAGCTTCTGATAATTGGAGGTGTTGCAGCC
CGGGATCAGCTCTCTGTTTTGGAAAATGGAGTAGATATAGTTGTAGGTACTCCGGGAAGACTAGATGACTTGGTG
TCAACTGGAAGAGCTGAAGTTATCTCAAGTTAGATTCTCGGTCCTGGATGAAGCTGATGGGCTTCTTTCTCAAGGT
TATTCTGATTTTATAAATAGGATGCACAATCAGATTCTCAGGTTACCTCTGATGGAAGAAAGACTTCAGGTGATT
GTTTGTCTGCTGCACTTTGCATTCTTTCGATGTAAAGAACTGTCCGAGAAGATAATGCATTTTCTACATGGGTT
GACTTAAAAGGAGAAGACTCTGTTCCAGATACTGTACACCATGTTGTTGTGCCAGTAAATCCCAAACTGACAGA
CTCTGGGAAAGGCTTGGAAAGAGCCACATTAGAAGTATGATGTACATGCAAAAGATAACACAAGACCTGGTGCT
AATAGTCCAGAGATGTGGTCTGAAGCTATTAAATCCTGAAAGGGGAGTATGCTGTCCGGGCAATCAAGGAACAT
AAGATGGATCAAGCAATTATCTTCTGTAGAACCAAAATTGACTGTGATAACTTGGAGCAGTACTTTATACAACAA
GGAGGAGGACCTGATAAAAAAGGACACCAGTTCTCATGTGTTTGTCTTCATGGTGACAGAAAGCCTCATGAGAGA
AAGCAAACTTGGAAAGATTTAAGAAAGGAGATGTAAGATTCTTGATTGACAGATGTAGCTGCTAGAGGAATT
GATATCCACGGTGTTCCTTATGTTATAAATGTCACTCTGCCGATGAAAAGCAAACTACGTACATCGAATTGGC
AGAGTAGGAAGAGCTGAAAGGATGGGTCTGGCAATTTCCCTGGTGGCAACAGAAAAAGAAAGGTTTGGTACCAT
GTATGTAGCAGCCGTGGAAAAGGGTGTATAACACAAGACTCAAGGAAGATGGAGGCTGTACCATATGGTACAAC
GAGATGCAGTTACTATCTGAGATAGAAGAACACCTGAACTGTACCATTCTCAGGTTGAGCCGGATATAAAGGTA
CCAGTGGATGAATTTGATGGGAAAGTTACCTACGGTCAGAAAAGGGCTGCTGGTGGTGAAGCTATAAAGGCCAT
GTGGATATTTTGGCACCTACTGTTCAAGAGTTGGCTGCCCTTGAAGAGGAGGCGCAGACATCTTCTGCTATCTT
GGCTACCTTCCTAACCAGCTGTTTCAGAACCTTCTGATTTTACATTTACTGAATAAGATTTGAGTAATGAAAGTC
TGAGTCTTAAACTCTAAACAGTTGTACTGCTTCCAAGCAGCAGTATTTATAGTAACGTAAGCTATTAATGCT
AACTCTTGATGTCAAGAAACATTAGTCTTAGGAATTCTTCAAAAAATGGCATCCCAATGAAAATAAATTTGATG
ACTATA

710/6881
FIGURE 663

MAAFSEMGVMPEIAQAVEEMDWLLPTDIAESIPLILGGGDVLMMAETGSGKTGAFSIPVIQIVYETLKDQQEGK
KGKTTIKTGASVLNKWQMNPHYDRGSAFAIGSDGLCCQSREVKWHGCRATKGLMKGKHYYEVSCHDQGLCRVGWS
TMQASLDLGTDKFEGFGGTGKKSHNKQFDNYGEEFTMHDTIGCYLDIDKGHVKFSKNGKDLGLAFEIPPHMKNQ
ALFPACVLKNAELKFNFGEFEFKFPPKDGFFVALSKAPDGYIVKSQHSGNAQVTQTKFLPNAPKALIVEPSRELAE
QTLNNIKQFKKYIDNPKLRELLIIGGVAARDQLSVLENGVDIVVGTPGRLLDLVSTGKLNLSQVRFLVLDEADGL
LSQGYSDFINRMHNQIPQVTS DGKRLQVIVCSATLHSFDVKKLSEKIMHFPTWVDLKGEDSVPD TVHHVVVPVNP
KTDRLWERLGKSHIRTD DVHAKDNTRPGANSPBMWSEAIKILKGEYAVRAIKEHKMDQAIIFCRTKIDCDNLEQY
FIQQGGGPDKKGHQFSCVCLHGDRKPKHERKQNLERFKKGDVRF LICTDVAARGIDIHGVPYVINVTLPDEKQNYV
HRIGRVGRAERMGLAISLVATEKEKVWYHVCSSRGKGCYNTRLKEDGGCTI WYNEMQLLSEIEEHLNCTISQVEP
DIKVPVDEFDGKVTYQGKRAAGGGSYKGHVDILAPT VQELAALEKEAQT SFLHLGYLPNQLFRTF

711/6881
FIGURE 664A

GAACATGGCGGCCCCCGAGTCAGGGCCGGCTTTGAGTCCAGGCACTGCAGAGGCCCTAGAGGCAACCAAAAAACATG
GTGCATCCTTTATCATCACGAAAGCAATTCGAGATCGTTTATTATTTTTACGCCAATACATCTGGTACAGCCCCG
CACCTTTTTTGTCTCCCTGATGGACTGGTTCGCTTGGTTAATAAACAGATAAACTGGCATTGTTGACTTGCAAGCA
ATAGCACTAGAGTGCATCTATACCTGTGAACGAAATGATCAACTCTGTCTTTGCTATGACCTACTAGAATGTCTG
CCAGAAAGAGGATATGGTGATAAGACAGAGGCAACCACAAAGCTTCATGACATGGTAGACCAACTGGAACAAATT
CTCAGTGTGTGACAGCTTTTGGAAAAACATGGACTCGAGAAACCAATTTCAATTTGTTAAAAACACTCAATCTAGC
TCAGAAGAGGCACGCAAGCTGATGGTTAGATTGACGAGGCACACTGGCCGGAAGCAGCCTCCTGTGCTGAGTCT
CATTGGAGAACGTTGCTGCAAGACATGTTAACTATGCAGCAGAATGTATACACATGCTAGATTCTGATGCCTGC
TATGAGATATTTACAGAAAGCCTTCTGTGCTCTAGTCGCCTTGAAAACATCCACCTGGCTGGACAGATGATGCAC
TGCAGTGCCTTGTTCAGAAAATCCTCCAGCTGGTATAGCCCATAAAGGGAAACCCCACTACAGGGTACAGTACGAA
AAGAGTATTGACTTGGTTTTGGCTGCCAGCAGAGAGTACTTCAATCTTCTACCAACCTCACTGATAGCTGCATG
GATCTAGCCAGGTGCTGCTTACAACCTGATAACAGACAGACCCCTGCCATTCAAGAGGAGCTAGATCTTATCCAA
GCCGTTGGATGTCTTGAAGAAATTTGGGGTAAAGATCCTGCCTTTGCAAGTGCGATTGTGCCCTGATCGGATCAGT
CTCATCAAGGAGTGTATTTCCAGTCCCCACATGCTATAAACAATCCACCAAGCTTCTGGGCCTTGCTGAGCTG
CTGAGGGTTGCAGGTGAGAACCAGAAAGAGCGGGGACAGGTTCTAATCCTTTTAGTGGAGCAGGCACTTCGC
TTCCATGACTACAAAGCAGCCAGTATGCATTGTGAGGAGCTGATGGCCACAGGTTATCCTAAAAGTTGGGATGTT
TGTAGCCAGTTAGGACAATCAGAAGGTTACCAGGACTTGGCCACTCGTCAAGAGCTCATGGCTTTTGCTTTGACA
CATTGCCCTCCTAGCAGCATTGAACCTCTTTTGGCAGCTAGCAGCTCTCTGCAGACAGAAATTCTTTATCAAAGA
GTGAATTTCCAGATCCATCATGAAGGAGGGGAAAATATCAGTGCTTACCATTAACTAGTAAAGCAGTACAAGAG
GATGAAGTAGGTGTTCCAGGTAGCAATTCAGCTGACCTATTGCGCTGGACCACTGCTACCACCATGAAAGTCCTT
TCCAACACCACAACCACCACCAAGCGGTGCTGCAGGCCGTGCTGATGGGCAGTGGTGGAAGAAGTCTTTAACT
TACCTTCGACCCCTTCAGGGGCAAAAATGTGGTGCTGCATATCAAATCGGAACCTACAGCCAATGAAGATCTAGAG
AAACAAGGGTGTCACTCTTTTATGAATCTGTCATCTCAAATCCTTTTGTGCTGAGTCTGAAGGGACCTATGAC
ACCTATCAGCATGTTCCAGTGGAAGCTTTGCAGAAGTATTGCTGAGAAGTGGAAAAATGGCAGAGGCTAAAAAT
AAAGGAGAAGTATTTCCAACAACTGAAGTTCTCTGCAACTAGCAAGTGAAGCCTTGCCAAATGACATGACCTTG
GCTCTTGCTTACCTTCTTGCTTACCACAAGTGTTAGATGCTAACCAGGTGCTTTGAAAAGCAGTCCCCCTCTGCA
TTATCTCTCAGCTGGCAGCGTATTACTATAGCCTCCAGATCTATGCCCGATTGGCCCCATGTTTCAGGGACAAG
TGCCATCTCTTTTACAGGCTGATCCCAAAGAATACTAAGATGGTCACCAGGCATGTGACTCGACATGAGCAGC
GAAGCCTGGCCTGAAGACCTTATTTCACTGACCAAGCAGTTACACTGCTACAATGAACGTCTCCTGGATTTCCT
CAGGCGCAGATCCTTCAGGGCCTTCGGAAGGGTGTGGACGTGCAGCGGTTTACTGCAGATGACCAGTATAAAAGG
GAAACTATCCTTGGTCTGGCAGAACTCTAGAGGAAAGCGTCTACAGCATTGCTATTCTCTGGCACACGTTAC
AGTGTCTCCCGCTGGGAAGTTTTTATGACCCATTGGAGTTTCTTTCACGGACAGTGGTTTGTCCACACTAGAA
ATTGAAAATAGAGCCCAAGACCTTCATCTCTTTGAGACTTTGAAGACTGATCCAGAAGCCTTTTACCAGCACATG
GTCAAGTATATTTACCCTACTATTGGTGGCTTTGATCACGAAAGGCTGCAGTATTATTTCACTCTTCTGGAAGAAC
TGTGGCTGTGCAGATTTGGGGAACTGTGCCATTAAACCAGAAACCCACATTGCACTGCTGAAGAAGTTTAAGGTT
GTTGCATCAGGTCTTAATTACAAAAAGCTGACAGATGAAAACATGAGTCTCTTGAAGCATTGGAGCCAGTTCTT
TCAAGTCAAAAATATCTTGTCTATTTCCAAACTTGTTCCTTCCAAAATCCCTGAAAAGGATGGACAGATGCTTTCCCA
AGCTCTCTGTACACCATCTGGTTACAGAAGTTGTTCTGGACTGGAGACCCTCATCTCATTAAACAAGTCCCAGGC
TCTTCACCGGAGTGGCTTCATGCCTATGATGTCTGCATGAAGTACTTTGATCGTCTCCACCCAGGTGACCTCATC
ACTGTGGTAGATGCAGTTACATTTTCTCCAAAAGCTGTGACCAAGCTGTCTGTGGAAGCCCGTAAAGAGATGACT
AGAAAGGCTATTAAGACAGTCAAAACATTTTATTGAGAAGCCAAGGAAAAGAACTCAGAAGACGAAGCTCAAGAA
GCTAAGGATTCTAAAGTTACCTATGCAGATACCTTGAATCATCTGGAGAAATCACTTGCCACCTGGAAACCTG
AGCCACAGCTTCATCCTTTCTCTGAAGAATAGTGAGCAGGAAACACTGCAAAAATACAGTCACCTCTATGATCTG
TCCCGATCAGAAAAAGAGAACTTCATGATGAAGCTGTGGCTATTTGTTTAGATGGTCAGCCTCTAGCAATGATT
CAGCAGCTGCTAGAGGTGGCAGTTGGCCCTCTTGACATCTACCCAAAGGATATAGTGCAGAGTGAATCATGAAA
ATAATTTCTGCATTGAGTGGTGGCAGTGTGACCTTGGTGGGCCAAGGGACCCACTGAAGGTCTGGAAGGTGTT
GTTGCAGCAGTCCACGCCAGTGTGGACAAGGGTGAGGAGCTGGTTTACCTGAGGACCTGCTGGAGTGGCTGCGG
CCTTTCTGTGCTGATGACGCTGGCCGGTGGGCCCGCATTCACGTGCTGCAGATTTGGGGCAATCATTTCAC

712/6881
FIGURE 664B

CTGACTGAGGAGGACAGCAAGCTCCTCGTGTTCTTTAGAACTGAAGCCATTCTCAAAGCCTCCTGGCCCCAGAGA
CAGGTAGACATAGCTGACATTGAGAATGAAGAGAACCGCTACTGTCTATTTCATGGAACTCCTGGAACTCTAGTCAC
CACGAGGCTGAATTTACGCACTTGGTTTTACTTTTGCAAGCTTGGCCACCTATGAAAAGTGAATATGTCATAACC
AATAATCCATGGGTGAGACTAGCTACAGTGATGCTAACCAGATGTACGATGGAGAACAAGGAAGGATTGGGGAAT
GAAGTTTTGAAAATGTGTCGCTCTTTGTATAACACCAAGCAGATGCTGCCTGCAGAGGGTGTGAAGGAGCTGTGT
CTGCTGCTGCTTAACCAGTCCCTCCTGCTTCCATCTCTGAAACTTCTCCTCGAGAGCCGAGATGAGCATCTGCAC
GAGATGGCACTGGAGCAAATCACGGCAGTCACTACGGTGAATGATTCCAATTGTGACCAAGAACTTCTTTCCCTG
CTCCTGGATGCCAAGCTGCTGGTGAAGTGTGTCTCCACTCCCTTCTATCCACGTATTGTTGACCACCTCTTGGCT
AGCCTCCAGCAAGGGCGCTGGGATGCAGAGGAGCTGGGCAGACACCTGCGGGAGGCCGCCATGAAGCCGAAGCC
GGGTCTCTCCTTCTGGCCGTGAGGGGGACTCACCAGGCCTTCAGAACCTTCAGTACAGCCCTCCGCGCAGCACAG
CACTGGGTGTGAGGGCCACCTGTGGCCCTGCTCCTTAGCAGAAAAAGCATCTGGAGTTGAATGCTGTTCCCAGAA
GCAACATGTGTATCTGCCGATTGTTCTCCATGGTTCCAACAAATTGCAAATAAACTGTATGGAACG

713/6881
FIGURE 665

ATGAGATCTCGGCTGCGAAGTGTGTCATCAGCATCTGGCTTTCTGGGTTATGATGTGGTCAGGGATGTGTCTTCT
GGCCCATTTTGGAGGCGCAAAATGGTGATCCATGTTTGGATTTATGTGGAAGGGACAGTTCCACAGTATGGAAA
AGGGGAGACAGTCCCAGGAAGCGTGAAGAGGAAAGGTCATTGGGAATCTCTCAGTACCTCTGCCCTAATCTCTTC
TCTCCTACGGAGGCAGTTAAGTCTAAAGCCATTTGTCTGGTTTCCATATCTGAACTTCAAGCAGCATCTTCTGGA
CTTAGTTCTTGGTGGCTGGAGCAGGGAAAGTGTGGCCTCATTGAAATGGGATCAGGTTTCAGGATGTTTCATGTG
TCTGTGTCTTTCTCACACATTGTGAGGAACAGGACAGTCCCTCTGATGCGCTGGTACCTGGCCGCCATGCAGATC
ATGAGGGGAGAGAAGCAGGCTGCTTCGGGGAATCCTGCCATAGAGGCTGGAGCTTCATGTGCATCTCAGCCAGCC
CTGCAGCAGCCCTCAGCAGCCCTTCTGTGGGGCTGGGGTATCTGTTGCCCAGGAAATCACTGGGCTGAGGAAGTA
GGAGATGAATGGCTCTGTGCTATCACACCATCTTCTGCATTGAGTCTTCTTCGCTTTCTTATCATCGTCATCATG
ATTATCATCAATAAATTTGTAAGGAAGGAAACAGGACTGCTTATTGATTTTCATTTCTATGCTACTGAGAGCCAA
GGCTACGGATTCTTTCCCCAGGAATGGAGAGAGGAAAGAGATGTGAGATTGGCTACCCAACTGTTGCATCAGTA
CCCCATTCTATCATCAACGGGTACAAACGAGTCCTGGCCTTGTCTGTGGAGACGGATTACACCTTCCCACTCGCT
GAAAAGGTCAAGGCCCTTCTTGGCTGATCCATCTGCCTTTGAGATGGAGGAGTGGAGGGGTGATGATAAAATGGAA
GCTCCCGTGTCAAGAGATCAGGCAAATGGGGAGACTCCGTGGGATCCTCTGAAGAATCTACATGCACGTACTTAC
GAGAATCAGATTAGGGATGCCTACCACACAGGAGTACCAATGTTTAGTATTAGCCCGTAA

714/6881
FIGURE 666

MRSRLRSVSSASGFLGYDVVRDVSSGPFWRRKNGDPCLDLCGRDSSTVWKRGDSPRKREEERSLGISQYLCPNLF
SPTEAVKSKAICLVSISELQAASSGLSSWWLEQGKCGLIEMSGFRMFHVSVSFSHIVRNRTVPLMRWYLAAMQI
MRGEKQAASGNPAIEAGWERTGAQM

715/6881
FIGURE 667

CGCGAAGAAGCTGGCAGGGGCACGAGCCGGGGCGGGTTTGAAGACGCGTCGTTGGGTTTTGGAGGCCGTGAAAC
AGCCGTTTGAGTTTGGCTGCGGGTGGAGAACGTTTGTACAGGGCCCGGCCAAGAAGGAGGCCCGCTGTTACGAT
GGTGTCCATGAGTTTCAAGCGGAACCGCAGTGACCGTTTCTACAGCACCCGGTGCTGCGGCTGTTGCCATGTCCG
CACCGGGACGATCATCCTGGGGACCTGGTACATGGTAGTAAACCTATTGATGGCAATTTTGCTGACTGTGGAAGT
GACTCATCCAACTCCATGCCAGCTGTCAACATTCAGTATGAAGTCATCGGTAATTACTATTCGTCTGAGAGAAT
GGCTGATAATGCCTGTGTTCTTTTGGCGTCTCTGTTCTTATGTTTATAATCAGTTCAATGCTGGTTTATGGAGC
AATTTCTTATCAAGTGGGTTGGCTGATTCCATTCTTCTGTTACCGACTTTTGGACTTCGTCTCAGTTGCCTGGT
TGCTATTAGTTCTCTCACCTATTTGCCAAGAATCAAAGAATATCTGGATCAACTACCTGATTTTCCCTACAAAGA
TGACCTCCTGGCCTTGGACTCCAGCTGCCTCCTGTTTCAATTGTTCTTGTGTTCTTTGCCTTATTCATCATTTTAA
GGCTTATCTAATTAACGTGTGTTTGGAACTGCTATAAATACATCAACAACCGAAACGTGCCGGAGATTGCTGTGTA
CCCTGCCCTTTGAAGCACCTCCTCAGTACGTTTGGCCAACCTATGAAATGGCCGTGAAAATGCCTGAAAAAGAACC
ACCACCTCCTTACTTACCTGCCTGAAGAAATTGTCCTTTGACAATAAATCCTATACCAGCTTTTGTGTTTAA
TGTTACAGAATGCTGCAATTCAGGGCTCTTCAAACCTGTTTATATAAATATGTTGCTTTTGTGTTAAGCATT
ATTTTCAAACACTAAGGAGCTTTTGGACATCTGTTAAACGTCCTTTTGTGTTTTTGTGTTAAGTCTTTTACATTTA
ATAGTTTTTGAAGACAATCTAGGTTAAGCAAGAGCAAAGTGCCATTGTTTGCCTTTAATTGGGGGTGGGAAGGG
AAAGAGGGTACTTGCCACATAGTTTCCCTTTTAACTGCACCTTCTTTATATAATCGTTTGCATTTTGTACTTGC
TACCTTGAGTACTTTTCAAGGAAGACTGACTTAAATATTGGGGTGAGTAAGTAGTTGGGTATAAGATCTGAACCTT
TCATCTGCAGAGGCAAGAAAAATATTTGACATTGTGACTTGACTGTGGAAGATGATGGTTGCATGTTTCTAGTTT
GTATATGTTTCCATCTTTGTGATAAGATGATTTAATAAATCTCTTTAAATACTT

716/6881
FIGURE 668

MVSMSEKRNRSDFYSTRCCGCCCHVRTGTIIILGTWYMVVNLLMAILLTVEVTHPNSMPAVNIQYEVIGNYYSSER
MADNACVLFAVSVMFIISSMLVYGAIISYQVGWLIPFFCYRLDFVLSCLVAISSLTYLPRKEYLDQLPDFPYK
DDLLALDSSCLLFIVLVFFALFIIFKAYLINCWNCYKYINNRRNVPEIAVYPAFEAPPQYVLPTYEMAVKMPKE
PPPPYLPA

717/6881
FIGURE 669

GCGGCTTGCGGGGACCACAGCTCCCGAAAGCGACGTTCCGGCCACCGGAGGAGCGGGAGCCAAGCAGGCGGAGCTC
GGCGGGAGAGGTGCGGGCCGAATCCGAGCCGAGCGGAGAGGAATCCGGCAGTAGAGAGCGGACTCCAGCCGGCGG
ACCCTGCAGCCCTCGCCTGGGACAGCGGCGCGCTGGGCAGGCGCCCAAGAGAGCATCGAGCAGCGGAACCCGCGA
AGCCGGCCCCGAGCCGCGACCCGCGCAGCCTGCCGCTCTCCCGCCGCCGGTCCGGGCAGCATGAGGCGCGCGGCG
CTCTGGCTCTGGCTGTGCGCGCTGGCGCTGAGCCTGCAGCCGGCCCTGCCGCAAATTGTGGCTACTAATTTGCCC
CCTGAAGATCAAGATGGCTCTGGGGATGACTCTGACAACTTCTCCGGCTCAGGTGCAGGTGCTTTGCAAGATATC
ACCTTGTCACAGCAGACCCCTCCACTTGGAAGGACACGCAGCTCCTGACGGCTATTCCCACGTCTCCAGAACCC
ACCGGCTTGAGGGCTACAGCTGCCCTCCACCTCCACCTGCCGGCTGGAGAGGGGGCCCAAGGAGGGAGAGGCTGTA
GTCCTGCCAGAAGTGGAGCCTGGCCTCACCGCCCGGGAGCAGGAGGCCACCCCGACCCAGGGAGACAGCATCA
GGGTTAAGAAGACTTTTTTTTTTTTTTTTAACTAGGAGAACCAAATCTGGAAGCCAAAATGTAGGCTTAGTTT
GTGTGTTGTCTCTTGAGTTTGTGCTCATGTGTGCAACAGGGTATGGACTATCTGTCTGGTGGCCCCGTTTCTGG
TGCTCTGTTGGCAGGCTGGCCAGTCCAGGCTGCCGTGGGGCCGCCGCTCTTTCAAGCAAGGCCAGAGGCCCCCA
GCCCAGGGCTCCTGCACTTACTTGCTTATTTGACAACGTTTCAGCGACTCCGTTGGCCACTCCGAGAGGTGGGCC
AGTCTGTGGATCAGAGATGCACCACCAAGCCAAGGGAACCTGTGTCCGGTATTCGATACTGCGACTTTCTGCCTG
GAGTGTATGACTGCACATGACTCGGGGGTGGGGAAAGGGTCCGGCTGACCATGCTCATCTGCTGGTCCGTGGGAC
GGTGCCCAAGCCAGAGGCTGGGTTTCAATTTGTGTAACGACAATAAACGGTACTTGTCATTTTCGGGC

718/6881
FIGURE 670

ACTGCGCGCCCCGCGCGGAGTCCCCGCGCGCGTCATGAGTCCCCGCGCGGTGCTCGTCACCTCCAGGCGACTTCA
GAATGCCACACTGGCCTCGACCTGACTGTGCCCCAGCACCAGGAGGTACGGGGCAAGATGATGTCTGGACACGT
GGAGTACCAGATCCTGGTGGTGACCCGTCTGGCTGCGTTCAAGTCGGCCAAGCACAGGCCCCAGGATGTCGTCCA
GTTCTTGGTCTCCAAAAGTACAGCGAGATTGAGGAGTTTTACCAGAACTGAGCAGTCGTTATGCAGCAGCCAG
CCTCCCCCACTACCCAGGAAGGTCTGTTTGTGGGGAGTCTGACATCCGGGAGAGGAGAGCCGTGTTCAATGA
GATCCTGCGCTGTGTCTCCAAGGATGCCGAGTTGGCAGGCAGCCAGAGCTGCTAGAGTTCTTAGGTACCAGATC
CCCAGGGGCTGCAGGGCTCACCAGCAGAGATTCTCTGTCTGGATGGCACAGACAGTCAGACAGGGAATGATGA
AGAGGCTTTGACTTTTTTGAGGAGCAAGACCAAGTGGCAGAAGAGGGTCCGCGCGTCCAGAGCCTGAAGGGCGA
GGATGCTGAGGAATCCTTGGAGGAGGAGGAGGCGCTGGACCTCTGGGCATTATGCGCTCCAAGAAGCCCAAGAA
ACATCGGTGTGAAGGGAAGGGACTGGGCCCTGCAGGGTCAGAACCTCCCCACCCCAAGGGGAGGCCAGGCAGAAG
CCTGGGTACACAGCACCAGAACTGCATGGTTCCATTTCTCCGGGGCTGTGGGGCCAAAGTAGAAGCCTGCGGGC
TGCGGGAGCGGCTCTACCCTAGGAGCCAGAGCCCAATGTGTCTTATTCCCCGTGGACATGAAGGGGAGGGAGGG
TGTGGGGATGCCTTGCCAACCAGAAGCCAGCCCCAAGGATGAAGCAAGACATGTGGGGCCGTAGCGAGGTGTCA
CATGGGGCAGGGAAGCTTCATGCCACGGGTCTGCCAGCCCCAGCACAGACCCAAACTGGGGCTGGGCCTCTAT
CCCTCCTCTGCCCTGTTCGCATAGTAAGAAGGAGTGACCGGTATCCTCCCTTCCCCTACCCCTAAGCTGTAGCC
TGGGTGACTGACTGGCCTGGGCCTGGGGTGGGGACGTCCCCAAGCCAAATTACTCCAGGGCCTCTGCTCCTCGTG
GCTGCCAGGGGCTGCAGGGTCTGGGTGGGTCTCCAGGAGAGGAATACTGAGTGGGAGATCGGCTGTCTGGAGT
GTTCTGATGCAAGTCTCTCTCCTGAGCCTCCTCTGATGCAAGCTCTAAAGGGAGAAGTCAGGCCCTGCCTCT
CCAGGGTATAGACGGCCCTGCTAGGCCCCAGTTCTTCTCCTTCCCCCTTTCCAGGAAAGGCCAGCCAGTCCA
TGGCCTTCTTGGGCCTCTGGGCACAGAGCCAAATGTTCTGTCATTGCAGCTCTCAGCAAACCTGGGTCATAGCTTCC
CCACAGCTCAGCCTGGGGCCTGGGCAGGGCTCCCAGCCTGCACCGCCTCCCACTGCCAGTGGGGCATGATTCTCT
CAGGCTTCTGCCCCGAGGCCTTCGTCTCTCAGGGTCTGGACTTGGTCAGTGGCCTTTTACCAGTGGAGCTGCC
TTCCAGGGAGAAGGAGCCGTGCGCCAGGGCAGGGCCGTGCCTTAGACTTCTCCCGACCCCAAGAGCGCTGGTA
CACAGGTCTAGGCACCACACAGTGCTTTGGAAATTCTCAGTGAATGATGTTTAATAAAGCAAAAAATGTC

719/6881
FIGURE 671

MQSPAVLVTSRRLQNAHTGLDLTVPQHQEVRGKMMSGHVEYQILVVTRLAAFKSAKHPEDVVQFLVSKKYSEIE
EFYQKLSSRYAAASLPPLPRKVLFGESDIRERRAVFNEILRCVSKDAELAGSPELLEFLGTRSPGAAGLTSRDS
SVLDGTDSTGNDEEAFDFFEEQDQVAEEGPPVQSLKGEDAEESLEEEELDPLGIMRSKKPKKHRCEGKGLGPA
GSEPPHPQGRPGRSLGHSTQNCMPFSPGLWGQSRSLRAAGAALT LGARAQCVLFPVDMKGREGVGMP CQPEAQP
QG

720/6881
FIGURE 672

ATGGGTCAAAGTCAGAGTGGTGGTCATGGTCCTGGAAGTGGCAAGAAGGATGACAAGGACAAAAAAGAAATAT
GAACCTTCTGTACCAACTAGAGTGAGGAAAAAGAAGAAAACAAAGGGACCAGATGCAGTCAGCAAACCTGCCA
CTGGTGACACCTCACACTCAGTGCCGGTTAAAATTACTGACATTAGAGAGAATTAAAGACTATCTTCTCATGGAG
GAAGAATTCATTAGAAATCAGGAACAGATGAAACCATTAGAAGAAAAGCAAAGGAGGAAAGATCAAAGTGGAT
GATCTGAGGGGGACCCCGATACAAGGATCTTCTGGAACTGGCTGCTCGGTCCTGCTCAACCACAAGGTGCATGC
CGTGATGGGGTGCTGATGGATGACACGGATCCCTGGTCACAGTGATGAAGATGGAAAAGACCCCCCAGGAGACC
TATGCCAATATTGGGGGGCTGGACAACCAAATTCAGGAAATAAAGGAAGCTGTGGAGTTTCTCTCACCCATCCT
GAATATTATGAAGAGATGGGTAAAAAGCCTCCTAAGGGGGTCATTCTCTATGGTCCACCTGGCACAGGCAAAACC
TTGTTATCCAAAGCAGTAGCAAACCAAACCTCAGCCACTTTCTTGAGAGTGGTTGGCTCTGAACCTTATTCAGAAG
TACCTAGTTGCTGAAGAGCATGAACTATCCATCATGTTTACTGATGAAATTGGAGCCATTGGGACAAAAAGATAT
GACTCAAATTCGGTGGCCGCATTGACAGGAAGATCAAGTTCTCCCTGCCTGATGAAAGGACTAAGAAGCGTATC
TTTCAGAATCACACAAGCAGGATGACACTGGCCGATGAAGTAACCCTGGACGACTTGATCATGGTTAAAGATGAC
CTCTCTGGTGCTGACATCAAGGCAATCTGTACAGAAGCTGGTCTGATGGCCTCAAGAGAACGTAGAAATGAAAGTA
ACGAATGAATCTTCAAAAAATATAAAGAAATGTTCTTTATAAGAAACAGGAAGGCACCCCTGAGGGGCTCTATC
TCTAGTGAACCACAGCTGCCATCAGGAAAATGTTTGGGCGATTCTCTGACCCCTGAAAAGGATGAGCAACTTGTT
CCCAAAGCTGGAGAAGACACTCCTGAGGGCTATTTAGGACAACCTTATGACTCAGCTCTTTGAGCAGAAAGAGGCC
AAAAAGTTCAGCAGAAAAAGCCCTGAACCTTTGGAAGAGCTGGCTTCAAGCCTGGCTTAG

721/6881
FIGURE 673

MGQSQSGGHGPGSGKKDDKDKKKKYEPSVPTVRKKKKKTKGPDVSKLPLVTPHTQCRLKLLTLERIKDYLLME
EEFIRNQEOMKPLEEKQKEERSKVDDLRGTPIQGSSGNWLLGPAQPQGACRDGVLMDDDTDLVTVMKMEKTPQET
YANIGGLDNQIQEIKEAVEFPLTHPEYYEEMGKKPPKGVILYGPPTGKTLLSKAVANQTSATFLRVVGSELIQK
YLVAEEHEL SIMFTDEIGAIGTKRYDSNSGGRIDRKIKFSLPDERTKKRIFQNHTSRMTLADEVTLDDLIMVKDD
LSGADIKAICTEAGLMASRERRMKVTNEFFKKYKEMFFIRNRKAPLRGSISSPEQLPSGKWLGDSSTPEKDEQLV
PKAGEDTPEGYLQQLMTQLFEQKEAKKFSRKSPELLEELASSLA

722/6881
FIGURE 674

GACGGCGCCTTCGCGAAACACTATGCTAATGGCATGGTGCCGCGGTCTGTCTTGCTGTGCCTGCGGCAGGGGCT
CGGAACCAATTTCATTCTGCACGGCCTGGGGCAGGAGCCCTTCGAGGGAGCTCGGTCACTGTGTTGCAGGTCCTC
GCCTAGAGACCTGCGAGATGGAGAAAGAGAGCACGAGGCGGCACAAAGGAAAGCCCCAGGAGCAGAGTCTTGCCC
ATCTCTCCCTCTGAGCATCTCGGACATTGGGACTGGATGTCTTTTCGTCACTGGAAAACCTCAGACTGCCGACGCT
GCGGGAAGAGTCATCACCTCGAGAGCTCGAGGACTCGAGCGGAGACCAGGGCCGGTCCGGTCCCACACACCAGGG
ATCCGAGGATCCTTCGATGCTCTCGCAGGCCAGTCCGCTATCGAGGTCGAAGAGCGTCACGTCTCCCTTCTTG
TTCAACTTCCAGAGAGAGACCCTTTCAGGCTGGGGAGCTGATTTTAGCTGAGACTGGGGAGGGAGAAACAAAATT
TAAGAAATTATTTAGGTTGAACAACTTCGGACTCTAAATAGTAACTGGGGGGCAGTCCCGTTCGGCAAGATCGT
GGGGAAGTTCCCCGCCAGATACTGAGGAGTTCCTTCGGTAAGCAGTACATGCTGAGGAGGCCAGCCTTGAAGA
CTATGTAGTATTGATGAAAAGAGGGACTGCCATAACATTCCCAAAGGATATTAATATGATTCTCTCAATGATGGA
TATCAACCCAGGTGATACTGTTTTGGAAGCTGGCTCAGGCTCTGGTGAATGAGCTTATTTTTATCCAAAGCAGT
TGGATCACAAGGACGAGTCATAAGTTTTGAGGTACGAAAAGACCACCATGATCTGGCTAAGAAGAATTACAAACA
CTGGCGTGATTCATGGAAATTAAGTCATGTAGAAGAGTGGCCAGACAATGTGGATTTTATTCATAAGGACATTC
AGGAGCAACCGAAGACATAAAATCTTTAACATTTGACGCAGTAGCTTTGGATATGTTAAATCCTCATGTTACTTT
GCCTGTTTTTTACCCACATCTTAAGCATGGTGGTGTATGTGCTGTATATGTAGTAAACATCACACAGGTTATTGA
ACTTTTAGATGGAATTGCGACCTGTGAACCTGCTCTTTCATGTGAAAAGATAAGCGAGGTCATTGTCAGAGATTG
GTTGGTTTGCCTTGCAAAACAGAAAAATGGAATTTTAGCTCAAAAAGTAGAATCTAAATCAACACAGATGTACA
ACTAGATTCTCAAGAGAAAATTGGAGTTAAAGGTGAGCTGTTTCAAGAGGATGACCATGAAGAATCGCATTCTGA
TTTTCCATATGGATCATTTCCTATGTTGCTAGACCAGTACACTGGCAACCTGGTCATACAGCTTTTTCTGTCAA
GTTGAGGAAGGTCAAACCACAACTTAACTGAGTACTCCAGATGACAGTAACTGACTTGAAGATGGAAAAATATCA
AAATAGAACTTTATATTGAAAATCACTGCTTCCATAGATTGGCATTTTTAGCTATTACTATGACTTATATACTT
ATACATATAATTTTGAAAATAACAACATAAAGATGTATAACATAGCAAACTGCTTAAACATCCCATTTTGACAC
TTGTCTTGCAAGTTAGTTTGACATTTTGTAGTTAATGATTCCAAATTGGTTTAGTTGGGCCATCTCATTCTTCACT
TCCTGTAAACCACTCCATAGATTTGTCTTTCTTCAAGAAATTAGTTTTCTTCTTTATTTGATTGATGGTCATT
GACTACTGAAATAAAATATGCATTTTAAGATAAAAAAAAAAAAAA

723/6881
FIGURE 675

MLMAWCRGPVLLCLRQGLGTNSFLHGLGQEPFEGARSLCCRSSPRDLRDGEREHEAAQRKAPGAESCPSLPLSIS
DIGTGCLSSLENLRLPTLREESSPRELEDSSGDQGRCPHQGSEDPMSLSQAQSAIEVEERHVSPSCSTSRERP
FQAGELILAETGEGETKFKKLFRLNNFGLLSNNGAVPFGKIVGKFPGQILRSSFGKQYMLRRPALEDYVVLTKR
GTAITFPKDINMILSMMDINPGDTVLEAGSGSGGMSLFLSKAVGSQGRVISFEVRKDHHDLAKKNYKHWRDSWKL
SHVEEWPDNVDFIHKDISGATEDIKSLTFDAVALDMLNPHVTLVPFYPHLKHGGVCAVYVNIQVIELLDGIRT
CELALSCEKISEVIVRDWLVLAKQKNGILAQKVESKINTDVQLDSQEKIGVKGELFQEDDHEESHSDFPYGSFP
YVARPVHWQPGHTAFLVKLRKVKPQLN

724/6881
FIGURE 676

ATGCAACAAGCTTCCTACCCCCAACCCGGCCAGACAAAGCCCCGGCAGCAGGGCCGGCACTTCCGAGCCTCCGGA
CTCGGGACGCCGCGCTCTTACCCTGGCCTCATCCAGAGTGACGCCGGCGCCACCGCTTCCCCGGCTTTGGCGGC
GGGGCAGCTGGCGGCGCGGGTCTTGGAGGAGCGGGTCCGAGAGGAGATGAAATGGCTGCTGCCTCCGGTCGCCCC
AGGCTCTGCTCCGGCCCCAGGCCCAGGCCCCGGGACCAGGAGACTTGGACCCGAGAAGGCGGAGAGAGCTCTTCCG
GGTGTTCACCATGGTCCAGCCAGGGGGACGGAGTCCACGCCGCGCCCGGGAGAGCCGAGCAAGGCCGGCCCCGCCG
GCCGCTGCCCCGCCCCCGGCGCGGCGTCAGCGCTCGGCGGCCGGGGAGCGACGCTCTAGGCTGTCAGCTCGGTGG
TTTCCAGCTCTCCCCGACCGCAGGGCTGGGGCGACCAGAAAAGCCAGACTAATCAGGACGAGTTCGGGCGTGGA
TTGGGCCTGAAGCCCTCATGCCCAGCCAGGCCTGAATGCCCCAGACCCACCGTCACCCCGTGGCCTGGATGGA
GCAGGGGACGGCCGGCGAAGGCGCGGGTGAAGGGGAGCGGCGACGAAGGCGCGAGGAGCTGACTGGGAATCCC
CAAGCTACCCGAGGCAAGACACTGAACTTGAGTCGCACTTTTCATGCCTGTAGAAATGAGAATGACAGCTGTGAT
AAAAAAAAAATTCATCTGGGCGGTATCTGGAACCTGGGCGTTCCGAACTTACAGTTATGCGCAACACAATGTTT
CTATTTTCAAAGGCGAACGCTGCACGATCGCAGTCCAGTGATACAAATAATAAAAGAGTTTATTACTTGTGAGCA
ATGATGAATGTTTCGTGTGGCATCCAGCGTTGAGTTCCTATTCTTACTTGGAACCTATTGGTCCCTGTGTCTGGGG
CCAGCACTGATGGACACAATAAGTTTGCACTCGTCTTCCCTCTCATGTATCATACCTGGAATGGGATCCAACAC
TTGATGTGGGACCTAGGAAAAGGCCTGAAGATTCCCCAGCTATACCAGTCTGGAGTGGTTGTCCTGGTTCTTACT
GTGTTATTGTAGGGCTGGCAGCTGTGTGA

725/6881

FIGURE 677

MQQASYPQPGQTKPRQQGRHFRASGLGTPRSYPGLIQSDAGATASPGFGGGAAGGAGLGGAGPRGDEMAAASGRP
RLCSGPRPRPGTRRLGPEKAERALPGVHHGPARGTESTPRPGEPSKAGPPAAAPPPARRQRSAAGERRSRLSARW
FPALPDRRAGATRKARLIRTSSGVELGLKPSCPARPECPRPHRHPRGLDGAGDGRRRRGVEGERRRRREELTGNP
QATRGKTLNLSRTFHACRNENDSCDKKKIHLGGIWNLGVPKLTVMRNTMFLFSKANAARSQSSDTNNKRVYYLSA
MMNVRVASSVEFLFLLGTHWSLCLGPALMDTTKFALVFPLMYHTWNGIQHLMWDLGKGLKIPQLYQSGVVVLVLT
VLFVGLAAV

726/6881
FIGURE 678A

ACAGAGAGATTGGTGTGTTTTGTGAGGCAGTGAGACCTAAGGTAACCTTTATCAAAAGGATGGAGTTGGGAAAAGG
AAACTACTCAGGACTGGACTGAATGCGTTGCATCAAGCAGTGCATCCGATCCATGGCCTTGCCCTGGACCGATGG
GAATCAAGTTGTCTTAAGTATTGCGGCTTCACAGTGGAGAGGTCAAGTTTGGGGACTCCAAAGTCATTGGACA
GTTTGAATGTGTCTGTGGGTTGTCTGGGCCCCACCTGTTGCAGATGATACACCTGTTCTACTCGCTGTCCAGCA
TGAGAAGCATGTCACTGTGTGGCAGCTGTGTCCCAGCCCTATGGAGTCAAGCAAATGGCTGACGTCTCAGACTTG
TGAGATTAGAGGATCACTACCTATCCTTCCCCAGGGCTGTGTGGCACCCTAAATGTGCTATTCTGACTGTGTT
GACTGCTCAGGATGTCTCCATTTTCCCTAATGTTCACTCTGATGATTCCCAGGTAAAGGCAGACATCAACACCCA
GGGCCGCATTCACTGTGCATGTTGGACCCAGGATGGCCTGAGGCTGGTGGTGGCAGTAGGCAGCAGCCTGCATT
TTATATTTGGGACAGCGCTCAGAAGACTCTTCACAGGTGCTCCTCCTGCCTGGTGTGTTGATGTGGACAGCCACGT
CTGCTCCATCACAGCAACTGTGGACTCACAGGTTGCTATAGCTACTGAGCTTCCATTGGATAAGATCTGTGGCTT
AAATGCATCTGAAACCTTTAATATCCCACCTAACAGTAAAGACATGACTCCGTATGCTTTACCAGTTATTGGTGA
AGTACGCTCTATGGATAAAGAGGCAACTGATTCTGAAACAAATCTGAAGTATCAGTTTCTTCTCTCTATTTAGA
ACCTCTGGATCTAACTCACATACATTTCAATCAACATAAGTCTGAGGGTAATTTCTTATTGTTCTAAGAAAAA
GGACTACTTGACAGGAAGTGGCCAAGATTCTTCACATTGGTCTTGTGACCTTAAAGAAGGCAGTTACCATGAC
GAGAAAAGTCACTATTCCAGGCATTCTGGTTCCTGATCTGATAGCATTAACTCTAAAGCCCACGTTAGTGGCAGT
GGCTTCCAACACTTGTAATATAATTTTGATCTACTCTGTCACTTCCATCTTCAGTCCCAAACATCCAGCAAATTCG
ATTAGAGAACTGAAAGACCAAAAGGGATATGTTTCTTGACAGACCAACTATTACTAATTTTGGTAGGAAAACA
AAAACCTCACTGATACAACTTTCTTCTCTTCAAAGTCTGATCAGTATGCCATTAGCTTGATTGTTAGAGAAAT
AATGTTGGAAGAAGAAGCTTCAATAACATCAGGTGAAAGCCAGACTACCTACTCTACTTTTCTAGTCTCCGTTAA
TAAAGCAAATAGAAAAAGTTAATTGAAAGTCTTTCCCCAGATTTTGTGACCAAAACAAAGGGCTGTGCTGAC
AGTTAATACCAGTAGTCAGAATGGAAGGCTGGAAGAAGTCTTATTAAAGAAATCCAGAGTCCCTCTGTCTAGTAT
CTGTGATGGCTCCATAGCTCTAGATGCTGAGCCTGTTACCCAGCCAGCATCGCTGCCAGACACAGCAGCACACC
AGACCACACAGCACATGAGCCTCCTCGTTTGCCTCAAAGAAAGAACTTACAAAGTGAAAGGAAAGTATATCA
GCTGTCTAAGGAGGTGGAATTTTATCTAGGAACCTGAGTTGAAATGCAACGGTGTCTTTCTGAACTTACAAACCG
TCTGCATAATGGGAAGAAATCCTCTCTCAGTGTATCCACTCTCTCAAGATCTTCTTATGTTTACATCATTATACCA
GAAACCTTATTATCTAGGTCTGTGTTGTTGAAAAAGAGCGGTGCTTCTCTGTGATGGTAAACTAAGGCTCAGTAC
AGTTTACGAGACTTTTGGCCTTTCTCTCAATTGAAATGCTACATGATTCCCACTGGATTCTTCTCTCTGTGACAG
TGAGGGCTTTATCCCGTTAACCTTACAGCCACACAGGAAATAATCATAAGAGATGGCAGCCTGTCCAGGTGAGA
TGCTTTCAGAGACTCTTTTCTCACAGTCCAGGTGCTGTTTCTTCTCTTAAAGTCTTTACAGGCTTGTGCTGCCCC
CAGTTTAGATACCACTGGCTGTTGTAACCATGTAGATGGCATGGCTTGATATCTGCAGTGTCTTGTCTGTGTAGC
TCTTCAGATGAGACCATTACAAACAAGGCTGCTTGACACTGGACACTCGCAATGAGACTCCCAGTGCCTCAG
GCGAAGCGCTTGCCATGGTCTGCTCTCTGTTTCCCCCTGTTTCCCCCTGAGCTGAGGCTCGCTGCTGTGTAGCA
GAGCTCAGTCTTTATTAGATGGCTCCGAAAGTGGTGTGTTATGTATTATGACTGTGTGGTTTTGACTAAGGGCAG
AATTCTCAGAACAAACAATATTATGGTGCCATATGGATGGTGTGTTTATGTTTCTCTGAGGCTTTGTGTCCCTT
GTCCAAAGCTGCATTGAAGCTGTCTTAGGAGCACTTAAAGATACTTGGCATTGTTATAGGTCTTTTCTTGGC
TTCAAGAGGAGGTTGAGGAGTCTGCTGGGGGCATGTGCTCTAGCATATTAACCTCAAACCAGCAAGAATTAGCA
GAGCTCCAAGGAGGACCAGAGACCCACTGGCTTCTGCTCTCAGGAACAGGAAGTGGCTCTGATGTTGCCTGGACC
TCCCAGAATTTAAACCAACCTCTTGTCTTCTTAAACAAATCTGGCTGACGAAGGTCCAGGTACTCTTAAAAAC
TGGCCCTGGGAAAATTTGAATGAAATTTCAAGGGAATTTGTCCCCTCTGGGTTCACCTTGAGGTTGTGCCGATG
CTGCTACCACACTGTGAGCCAGGTAAGTCTACTGCAGGATTTTGTGCTGTGGCCACTCATGAGTGTCCCTGA
AATAACTTTTTTTTTTTTTTAAATCCAGTTTTGGGATCACGCAACTTTCCTATTTTTCTCCAGTAGTCAGCTCCC
TTAGTTAACTTGTCATTTAATTTGATATTTTTATTTTCTCTCTTTTAAAGTCTTAGAGACCAGCAGAGAATCTG
TGAGAGAAAGTATTTAGGAAGTTAGAAATTCAACCGAATCTGAGGTAGTCTTAAAGTGGCAATTTTGTTCAC
TTATGGGCTAAAGTACCAGCTTAGTCAGGTAAGAGCCCTGACCCACTTCAGATGGTAACACCACCTTCTACTGCC
TTCAGATGGAATCACAGATTTTCACTACGGCGCATACAAATTTGATCAGTGAAGTGGCTAGGCATCTGCAGATAAA
TTGTTTTCAGCCATAGAAGCTCCATTAGCACATATGCTTCTTTTCCCCCTTCTTTTAAATCATCTGGAAGAA
ACTATTTTGTGCCCTTGGGACTCCTGTCTGTCTGTTACAGTTTACCAAGATGGAGCTGGGTTAGGAAAGAGTG
AGGGCCCATTTTGTGGTTCAAGTGCCTAGACAGCTGCTGGGGTAGGAAGCACAGGCAATGTCTGCAATCAGCTG

727/6881

FIGURE 678B

TGGGAGAGCGGTGACTGAGAACAGTCTGAGGCCTGGCTCCACTTGAAGTATCTGGGGTGCGATGAAATCACAAT
TATCTTGAAGCCTAAAGAGGGAACTACAAGACTGTAACTAAGATCAATGTGGGCACCTAAAAGGGTATGTTAAA
ATCACCATTCTCAGGTCAAATACTGTGAAATAAAATGGAAACGGAAAAAAAAAAAAAAAAAAAAAAAAAAAAA
AAAAAAAAA

728/6881

FIGURE 679

MELGKGKLLRTGLNALHQAVHPIHGLAWTDGNQVVLTDLRLHSGEVKFGDSKVIQFECVCGLSWAPPVADDTVP
LLAVQHEKHVTVWQLCPSPMESSKWLTSTCEIRGSLPILPQGCVWHPKCAILTVLTAQDVSIFPNVHSDDSQVK
ADINTQGRIHCACWTQDGLRLVVAVGSSLHSYIWDSAQKTLHRCSSCLVFDVDVSHVCSITATVDSQVAIATELPL
DKICGLNASETFNIPPNSKDMTPYALPVI GEVRSMDKEATDSETNSEVSVSSSYLEPLDLTHIHFNQHKSEGNL
ICLRKKDYLTGTGQDSSHLVLTFFKKAVTMTRKVTIPGILVPDLIAFNLKAHVAVASNTCNIIIIYSVIPSSVP
NIQQIRLENTERPKGICFLTQDQLLLILVGKQKLTDTTFLPSSKSDQYAI SLVREIMLEEEPSITSGESQTTYST
FSAPLNKANRKKLIESLSPDFCHQNKGLLLTVNTSSQNGRPGRTLKEIQSPLSSICDGSIALDAEPVTQPASLP
RHSSTPDHTSTLEPPRLPQRKNLQSEKETYQLSKEVEILSRNLVEMQRCLSEL TNRLHNGKKSSSVYPLSQDLPY
VHIYQKPYLGPVVEKRAVLLCDGKLRRLSTVQQTGFLSLIEMLDHSHWILLSADSEGF IPLTFTATQEI IIRDG
SLSRSDVFRDSFSHSPGAVSSLKVFTGLAAPSLDTTGCCNHVDGMA

729/6881

FIGURE 680

TTCTCTTAGCAGCACCCAGCTTGCCACCCATGCTCAAGATGGGCGGGATGCCAGCCTGTTACATAAAATGTGCCA
AAAGCCTGGCCATGCCTGGAAAATGGACCAATCCGCCCCCAAGAGGTGGGTCTCGTTCCCTAGAGAGAAGGAA
GTTTCCTCTCCTTGAAGTGAGAGCTAGAATCGCACTTTCTGTCAAGCTGAGAGAAAAGACTCTTTTCCAGAGGCTA
AAAGGACAAGAAAATCTGATTTGCTTGCTTCTAACTTTGCGTTTTAAAGGGGGAAGGAGGAAAGGAAAGAGGGGG
AGGGTGGTTCTGCTTAGCCCCACCCCTCCGGCTACCCAGGTCCAGCCGTCCATTCCGGTGGAGGCAGAGGCAGT
CCTGGGGCTCTGGGGCTCGGGCTTTGTACCGGGACCCGAGGAGCCAGAACCACTCGGCGCCGCCCTGGTGATG
GGAGGGGAGCCGGGCCAGGAACAATATGTTAGCCGTGCACTTTGACAAGCCGGGAGGACCGGAAAACCTCTACGT
GAAGGAGGTGGCCAAGCCGAGCCCGGGGAGGGTGAAGTCCTCCTGAAGGTGGCGGCCAGCGCCCTGAACCGGGC
GGACTTAATGCAGAGACAAGGCCAGTATGACCCACCTCCAGGAGCCAGCAACATTTTGGGACTTGAGGCATCTGG
ACATGTGGCAGAGCTGGGGCCTGGCTGCCAGGGACACTGGAAGATCGGGGACACAGCCATGGCTCTGCTCCCCGG
TGGGGGCCAGGCTCAGTACGTCACTGTCCCCGAAGGGCTCCTCATGCCTATCCCAGAGGGATTGACCCTGACCCA
GGCTGCAGCCATCCCAGAGGCCTGGCTCACCGCCTTCCAGCTGTTACATCTTGTGGGAAATGTTGAGGCTGGAGA
CTATGTGCTAATCCATGCAGGACTGAGTGGTGTGGGCACAGCTGCTATCCAACCTACCCGGATGGCTGGAGCTAT
TCCTCTGGTCACAGCTGGCTCCCAGAAGAAGCTTCAAATGGCAGAAAAGCTTGAGCAGCTGCTGGATTCAATTA
CAAAAAGAGGATTTCTCTGAAGCAACGCTGAAATTCACCAAAGGTGCTGGAGTTAATCTTATTCTAGACTGCAT
AGGCGGATCCTACTGGGAGAAGAACGTCAACTGCCTGGCTCTTGATGGTCGATGGGTCTCTATGGTCTGATGGG
AGGAGGTGACATCAATGGGCCCTGTTTTCAAAGCTACTTTTTAAGCGAGGAAGTCTGATCACCAGTTTGCTGAG
GTCTAGGGACAATAAGTACAAGCAAATGCTGGTGAATGCTTTCACGGAGCAAATCTGCCTCACTTCTCCACGGA
GGGCCCCAACGTCTGCTGCCGGTCTTGACAGAATCTACCCAGTGACCGAAATCCAGGAGGCCATAAGTACAT
GGAGGCCAACAAGAACATAGGCAAGATCGTCTGGAAGTGGCCAGTGAAGGAGGATGGGGCAGGACAGGACGCG
GCCACCCAGGCCTTTCCAGAGCAAACCTGGAGAAGATTACAATAGACAGGCCAAGAAACCCGGTGCTTCCTCC
AGAGCCGTTTAAAGCTGATATGAGGAAATAAAGAGTGAAGTGG

730/6881
FIGURE 681

GGCACGAGGGTCCCCTTACCGCCTCCTGCTCCTGCCGCGCGCACCCCCGGGGCTTCGGCTCCGGCACGGGTTCGCG
CCCAGCTTTCCTGCACCTGAGGCCGCCGCCAGCCGCCGCCATGGGTGCCTACCTCTCCCAGCCCAACACGGTGA
AGTGCTCCGGGGACGGGGTCGGCGCCCCCGCCTGCCGCTGCCCTACGGCTTCTCCGCCATGCAAGGCTGGCGCG
TCTCCATGGAGGATGCTCACAACGTATTCTGAGCTGGACAGTGAGACAGCCATGTTTTCTGTCTACGATGGAC
ATGGAGGGGAGGAAGTTGCCTTGTACTGTGCCAAATATCTTCTGATATCATCAAAGATCAGAAGGCCTACAAGG
AAGGCAAGCTACAGAAGGCTTTAGAAGATGCCTTCTTGGCTATTGACGCCAAATTGACCACTGAAGAAGTCATTA
AAGAGCTGGCACAGATTGCAGGGCGACCCACTGAGGATGAAGATGAAAAAGAAAAAGTAGCTGATGAAGATGATG
TGGACAATGAGGAGGCTGCACTGCTGCATGAAGAGGCTACCATGACTATTGAAGAGCTGCTGACACGCTACGGGC
AGAACTGTCAAGGGCCCTCCCCACAGCAAATCTGGAGGTGGGACAGGCGAGGAACAGGGTCCCAGGGCCTCA
ATGGGGAGGCAGGACCTGAGGACTCAACTAGGGAACTCCTTCACAAGAAATGGCCCCACAGCCAAGGCCTACA
CAGGCTTTTCTCCTCAACTCGGAACGTGGGACTGAGGCAGGCCAAGTTGGTGAGCCTGGCATTCCCCTGGTGAGG
CTGGGCCTTCTGCTCTTCAGCCTCTGACAAGCTGCCTCGAGTTGCTAAGTCCAAGTTCTTTGAGGACAGTGAGG
ATGAGTCAGATGAGGCGGAGGAAGAAGAGGAAGACAGTGAGGAATGCAGCGAGGAAGAGGATGGCTACAGCAGTG
AGGAGGCAGAGAATGAGGAAGATGAGGATGACACCGAGGAGGCTGAAGAGGACGATGAAGAAGAAGAAGAAGAGA
TGATGGTGCCAGGGATGGAAGGCAAAGAGGAGCCTGGCTCTGACAGTGGTACAACAGCGGTGGTGGCCCTGATAC
GAGGGAAGCAGTTGATTGTAGCCAACGCAGGAGACTCTCGCTGTGTGGTATCTGAGGCTGGCAAAGCTTTAGACA
TGTCTATGATCACAAACCAGAGGATGAAGTAGAACTAGCACGCATCAAGAATGCTGGTGGCAAGGTCACCATGG
ATGGGCGAGTCAACGGGGGCCTCAACCTCTCCAGAGCCATTGGGGACCACTTCTATAAGAGAAACAAGAACCTGC
CACCTGAGGAACAGATGATTTAGCCCTTCTGACATCAAGGTGCTGACTCTCACTGACGACCATGAATTCATGG
TCATTGCCTGTGATGGCATCTGGAATGTGATGAGCAGCCAGGAAGTTGTAGATTTCATTCAATCAAAGATCAGCC
AGCGTGATGAAAATGGGGAGCTTCGGTTATTGTATCCATTGTGGAAGAGCTGCTGGATCAGTGCCTGGCACCAG
ACACTTCTGGGGATGGTACAGGGTGTGACAACATGACCTGCATCATCATTTGCTTCAAGCCCCGAAACACAGCAG
AGCTCCAGCCAGAGAGTGGAAGCGAAACTAGAGGAGGTGCTCTCTACTGAGGGGGCTGAAGAAAATGGCAACA
GCGACAAGAAGAAGAAGGCCAAGCGAGACTAGCAGTCATCCAGACCCCTGCCACCTAGACTGTTTTCTGAGCCC
TCCGGACCTGAGACTGAGTTTTGTCTTTTTCTTTAGCCTTAGCAGTGGGTATGAGGTGTGCAGGGGGAGCTGGG
TGGCTTCACTCCGCCCATTCCAAAGAGGGCTCTCCCTCCACACTGCAGCCGGGAGCCTCTGCTGTCTTCCCAGC
CGCTCTGCTCCTCGGGCTCATCACCAGTTCTGTGCCTGTGCTCTGTTGTGTGGAGGGAAGGACTGGCGGTTCT
GGTTTTTACTCTGTGAACTTTATTTAAGGACATTCTTTTTTATTGGCGGCTCCATGGCCCTCGGCCGCTTGCAAC
CGCTCTCTGTTGTACACTTTCAATCAACACTTTTTTCACTAAAGGCCAAAACCTAAAAAAAAAAAAA

731/6881

FIGURE 682

MGAYLSQPNTVKCSGDGVGAPRLPLPYGFSAMQGWRVSMEDAHNCIPELDSETAMFSVYDGHGGEEVALYCAKYL
PDI IKDQKAYKEGKLQKALEDAFLAIDAKLTTEEVIKELAQIAGRPTEDEDEKEKVADEDDVDNEEAALLHEEAT
MTIEELLTRYGQONCHKGPPHSKSGGGTGEEPGSQGLNGEAGPEDSTRETSPQENGPTAKAYTGFSNSERGT
EAGQVGEFGIPTGEAGPSCSSASDKLPRVAKSKFFEDSEDESDEAEEDSEECSEEDGYSSEEAENEDEDDTEE
AEEDDEEEEEEEMVPGMEGKEEFGSDSGTTAVVALIRGKQLIVANAGDSRCVSEAGKALDMSYDHKPEDEVELA
RIKNAGGKVTMDGRVNGGLNLSRAIGDHFYKRKNLPPEEQMISALPDIKVLTLLTDDHEFMVIACDGIWNVMSSQ
EVVDFIQSKISQDENGELRLSSIVEELLDQCLAPDTSGDGTGCDNMTCIICFKPRNTAELQPESGKRKLEEV
LSTEGAEENGNSDKKKKAKRD

732/6881

FIGURE 683

CCACTGCCGCTGCCGCCGCCCTCTCCTTAGTCGCCGGCATGACGACCGCGTCTACCTCGCAGGTGCGCCAGAAGT
GCTGAGAAATTGATGAAGCTGTAGAACCAACGAGGTGGCCGAATCTTCCTTCAGGATATCAAGAAACCAGACTGT
GCGGGGAGAATGCGATGGGAGAGCGGGCTGAATGCGATGGATTACATTTGAAAAAATTGTGCATTTGCATTAC
ATTTGAAAAAATGTGAATCAGTCACTACTGGAAGTGCACAACTGGCCACTGACAAAAATGACCCCCATTTGT
GTGACTTCATTGAGACACATTACCTGAATGAACAAGTGAAGGCCATCAAAGAATTGGGTGACCACGTGACCAACA
TGCACGAGATGGGAGCGCCCGAATCTGGCGTGGCAGAATACCTCTTTGACAAGCACACCCTGGGAGACAGTGATA
ATGAAAGCTAAGCCTCAGGCTAATTTCCCATAGCCATAGGGTGAATTACCTTGTACCAAGGCAGCGCATGTAT
GTGGGGTTTCCTTTACCTTTTCTATAAGTTGTTCCAAGACACCCACTTAAGTTCTTTGATTGTACCATTCTT
CAAATAAATAAATTTGGTACCC

733/6881

FIGURE 684

CCCTATCCGGACAGGTGGCTCTTGCCCTTTAGACTACAGTTCCCAGCATGCCAGGCGATTGCGTCCCAGAACCG
ACGTCCCACCGCCTTCCCACATCGGATCGCAGGGCTCCAAAATGGCGAGTGAGGCTGCGGGGACTCGTGAGCA
GCGGAGGGGGAGCGTGCAGAGCCGCTGCGGCCCTCACAGTCCGGAGCCCGGCCGTGCCGTGCCGTAGGGAACATG

CACCTTTTCCATTCCCGAAACCGAGTCCCGCAGCGGGGACAGCGCGCGCTCCGCCTACGTGGCCCTATAACATTACAC
GTGAATGGAGTCTCTGCACTGTCGGGTGCGCTACAGCCAGCTCCTGGGGCTGCACGAGCAGCTTCGGAAGGAGTAT
GGGGCCAATGTGCTTCTCTGCATTCCCCCAAAGAAGCTTTTCTCTGTGACTCCTGTGAGGTAGAACAGAGGAGA
GAGCAGTTAGAGAAGTACATGCAAGCTGTTTCGGCAAGACCCATTGCTTGGGAGCAGCGAGACTTTCAACAGTTC
CTGCGTCGGGCACAACAGGAGACACAGCAGGTCCCCACAGAGGAAGTGCTTGGAAAGTGCTGCTCAGCAACGGG
CAGAAAGTTCTGGTCAACGTGCTAACTTCAGATCAGACTGAGGATGTCTTGGAGGCTGTAGCTGCAAGCTGGAT
CTTCCAGATGACTTGATTGGATACTTTAGTCTATTCTTAGTTCGAGAAAAAGAGGATGGAGCCTTTTCTTTTGTA
CGGAAGTTGCAAGAGTTTGAGCTGCCTTATGTGTCTGTACCAGCCTTCGGAGTCAAGAGTATAAGATTGTGCTA
AGGAAGAGTTATTGGGACTCTGCCTATGATGACGATGTCATGGAGAACCGGGTTGGCCTGAACCTGCTTTATGCT
CAGACGGTATCAGATATTGAGCGTGGGTGGATCTTGGTACCAAGGAACAGCACCGGCAACTCAAATCTCTGCAA
GAGAAAGTCTCCAAGAAGGAGTTCTTGAGACTGGCCCAGACGCTGCGGCACTATGGCTACTTGCGCTTTGATGCC
TGTGTGGCTGACTTCCCAGAAAAGGACTGTCCTGTGGTGTGTGAGCGCGGGCAACAGTGAGCTCAGCCTGCAGCTC
CGCCTGCCTGGCCAGCAACTCCGAGAAGGCTCCTTCCGGGTACCCCGCATGCGATGCTGGCGGGTCACTTCCCTCT
GTACCATTGCCCAGTGGAAGCACGAGCAGCCAGGCCGGGGCGGGGTGAGGTGCGCCTGGAAGTGGCTTTTGAA
TACCTCATGAGCAAGGACCGGCTACAGTGGGTACCATCACTAGCCCCCAGGCTATCATGATGAGCATCTGCTTG
CAGTCCATGGTTGATGAACTGATGGTGAAGAAATCTGGCGGCAGTATCAGGAAGATGCTGCGCCGGCGGGTGGGG
GGTACTCTGAGACGCTCAGACAGCCAGCAAGCAGTGAAGTCCCCACCACTGCTTGAGTCACCTGATGCCACCCGG
GAGTCTATGGTCAAACCTCTCAAGTAAGCTGAGTGCCGTGAGCTTGCGGGGAATTGGCAGTCCCAGCACAGATGCC
AGTGCCAGTGATGTCCACGGCAATTTGCGCTTCGAGGGCATTGGAGATGAGGATCTGTAATCTCCACTGCTTGGA
TGTCTGCCCTCTACCCCAGAGGAATTTACAGAACTTGCCCTGTGCCTGTGTCCCCATGCTAGGGGCGGAGGGG
TCTTTTCTTCTTCTTCTTCTACCTACCCCTTTTCTCTTGGCCAGGGGCTCGTATCTACCTTCTTGTGCCCT
GGGCTGGCTGCACAGGAGATTGCCCTTCTTTTTCAGAGCTGGCCCTCGATGCCAAATTAGCATTTAGTATTTT
GCACAAAGTCTAAGGACCATGGCTGCCTGCCTTGGGGAGGAACCATAGCTCCCTCTGGGCCGCTTCTGGCCTCT
TGGAGCCATGGGCCAAAGGCCAAGGGGATGGGCAGAGGTCTGTGTTTGGTCTGGCCCACTTCCCCATCATTAAC
TCAGCCTGACTGCTGCCT

734/6881

FIGURE 685

MHFSIPETESRSGDSGG SAYVAYNIHVNGVLHCRVRYSQLLGLHEQLRKEYGANVLP AFPPKKLFS LTPAEVEQR
REQLEKYMQAVRQDPLLGSSETFNSFLRRAQQETQQVPT EEVSLEVLLSNGQKVLVNVLTSDQTEDVLEAVA AKL
DLPDDLIGYFSLFLVREKEDGAFSFVRKLQEFELPYVSVTSLRSQ EYKIVLRKSYWDSAYDDDV MENRVGLNLLY
AQTVSDIERGWILVTKEQHRQLKSLQEKVSKKEFLRLAQT LRHYGYLRFDACVADFPEKDCPVVVSAGNSELSLQ
LRLPGQQLREGSFRVTRMRCWRVTSSVPLPSGSTSSPGRGRGEVRLELAF EYLMKDR LQWVTITSPQAIMMSIC
LQSMVDELMVKKSGGSIRKMLRRRVGGTLRRSDSQAVKSPPLESPDATRES MVKLSSKL SAVSLRGIGSPSTD
ASASDVHGNFAFEGIGDEDL

735/6881
FIGURE 686

GATGGCTGCTGTGGCCGTGGCTGTTGCGGAGGACTCGGGATCCGGGATGAAGGCGGAGCTTCCCCCTGGGCCTGG
GGCAGTGGGGAGGGAAATGACCAAAGAAGAAAAGCTGCAGCTTCGGAAGGAAAAGAAACAGCAGAAGAAGAAACG
GAAGGAAGAAAAGGGGGCAGAACCCAGAGACTGGCTCTGCTGTATCTGCAGCCCAATGTCAAGTAGGCCCAACCAG
AGAACTGCCAGAATCGGGCATTGAGTTGGGCACTCCTCGGGAGAAAAGTTCCAGCTGGTCGGAGTAAGGCCGAAC
TCGGGCTGAGCGTCGAGCCAAGCAGGAGGCCGAGCGGGCCCTGAAACAGGCAAGAAAAGGGGAACAAGGAGGACC
ACCTCCTAAGGCCAGCCCCAGCACAGCTGGAGAAAACCCCTCAGGAGTGAAGCGTCTCCCTGAGTACCCTCAGGT
TGATGACCTACTTCTGAGAAGGCTTGTTAAAAAACCAGAGCGTCAACAGGTTCTTACACGAAAGGATTATGGATC
CAAAGTCAGTCTCTTCTCTCACCTACCCAGTACAGCAGACAAAACCTCTTGACCCAGTTTATGAGCATCCCATC
CTCTGTGATCCACCCAGCCATGGTGCAGTCTCGGCCTGCAGTACTCCAGGGCCTGGTCAGTGGCTCCAATGCCCC
GTGTATTGCCCTGCTTCGTGCCTTGACAGCAGGTGATTGAGGATTACACAACACCGCCTAATGAAGAACTCTCCAG
GGATCTAGTGAATAAACTAAAACCCCTACATGAGCTTCCTGACTCAGTGCCGTCCCCTGTGAGCGAGCATGCACAA
CGCCATCAAGTTCCTTAACAAGGAAATCACCAGTGTGGGCAGTTCCAAGCGGGAAGAGGAGGCCAAGTCAGAACT
TCGAGCAGCCATTGATCGGTATGTGCAAGAGAAGATTGTGCTAGCAGCTCAGGCAATTTACGCTTTGCTTACCA
GAAGATCAGTAATGGAGATGTGATCCTGGTATATGGATGCTCATCTCTGGTATCACGAATTTCTCAGGAGGCTTG
GACAGAGGGCCGGCGGTTTCGGGTGGTAGTGGTGGACAGCCGGCCATGGCTGGAAGGAAGGCACACACTACGTTT
TCTAGTCCATGCTGGTGTCCCAGCCTCCTACCTGCTGATTCCTGCAGCCTCCTATGTGCTCCCAGAGGTTTCCAA
GGTGCTATTGGGAGCTCATGCACTCTTGCCCAACGGGTCTGTGATGTACGCGGTAGGGACAGCACAGTTAGCCCT
GGTGGCTCGAGCCCATATGTACCAAGTGTGGTTTGCTGTGAAACATACAAGTTCTGTGAGCGTGTGCAGACTGA
TGCCTTTGTCTCTAATGAGCTAGATGACCCTGATGATCTGCAATGTAAGCGGGGAGAACATGTTGCGCTGGCTAA
CTGGCAGAACCACGCATCCCTACGGTTGTTGAATCTAGTCTATGATGTGACTCCCCCAGAGCTTGTGGATCTGGT
GATCACGGAGCTGGGGATGATCCCTTGCACTTCTGTACCTGTTGTTCTACGAGTCAAGAGCAGTGACCAGTGACG
GGGGAACACAGGGTTAATAAATGCCATACTCCCTACCCCTCAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA
AAAAAAAAAC

736/6881
FIGURE 687

MAAVAVAVREDSGSGMKAELPPGPGAVGREMTKEEKLQLRKEKKQKKKRKEEKGAEPETGSAVSAAQCQVGPTR
ELPESGIQLGTPREKVPAGRSKAELRAERRAKQEAERALKQARKGEQGGPPPKASPSTAGETPSGVKRLPEYPQV
DDLLLRRLVKKPERQQVPTRKDYGSKVSLFSLPQYSRQNSLTQFMSIPSSVIHPAMVRLGLQYSQGLVSGSNAR
CIALLRALQQVIQDYTTTPNEELSRDLVNKLKPYMSFLTQCRPLSASMHNAIKFLNKEITSVGSSKREEEAKSEL
RAAIDRYVQEKIVLAAQAISRFAYQKISNGDVILVYGCSLVSRLQEAWTEGRRFRVVVDSRPWLEGRHTLRS
LVHAGVPASYLLIPAASYVLPEVSKVLLGAHALLANGSVMSRVGTAQLALVARAHNPVLVCCETYKFCERVQTD
AFVSNELDDPDDLQCKRGEHVALANWQNHASLRLNLVYDVTPELVDLVITELGMIPCSSVPVVLRVKSSDQ

737/6881
FIGURE 688

AGGACTTTGGCGAGGGGGCAGCCATTTTGGGGGGTGTGATGGATACCTGCGGGGTCGGCTATGTTGCCCTGGGG
GAGGCCGGCCCCGTGGGGAACATGACTGTGGTAGACTCTCCTGGACAAGAGGTGCTAAATCAGCTTGATGTCAAG
ACCTCTTCAGAAATGACCAGTGCAGAGGCTTCCGTAGAGATGTCAATACCTACCCCTTGCCTGGATTGAGGAT
TCTCCTGATCAGAGGAGGCTCCCTCCAGAGCAGGAAAGCCTCTCCAGACTGGAACAGCCAGATCTTTCTTCAGAG
ATGTCAAAGGTCTCAAAGCCTAGGGCCTCAAAGCCTGGCCGGAAGAGAGGTGGTAGGACACGAAAAGGCCCCAAA
AGGCCCCAACAGCCTAATCCTCCATCAGCCCCACTGGTTCCTGGTCTCTTAGATCAATCCAACCCTCTGTCCACC
CCCATGCTAAGAAACGAGGTGAAAGTCCAAGGCAGAGCTGCTGCTGCTGAAGTTGTCAAAAGACCTAGATCGG
CCAGAATCTCAATCTCAAAGAGGCCCCCTGAGGACTTTGAGACCCCTTCTGGGGAACGACCCCGCGAAGGGCT
GCCAAGTGGCACTTCTGTATCTTCAGGAACGGCTGAAGAGCTCTAACAGCCCTGCCTGCCCCCTGTGTCTGT
CCTGAGGGCCCCAAGGTGAGCAGCCCCACCAACCGAAGAAGATCCGGCAGCCAGCAGCCTGTCCAGGTGGAGAA
GAGGTGGATGGTGTCTCCACGGGATGAAGACTTTTTTCTCCAGGTTGAGGCTGAAGATGTGGAAGAAAGTGAGGGC
CCAAGTGAGAGCTCATCTGAACCTGAGCCTGTAGTGCCCCGAAGCAGCCACGAGGATCTACTTCAGGGAACAG
AAACCACACTGCCGAGGAATGGCTCCCAATGGCTTACCAATCATATCATGGCTCCTGTTTGAAGTGCCTCCAT
CTCACCAGGACTTCCGAGAGCAGAAACATTCTACTGGGAGTTTGTGAGTGGATTCTTTAGCCTGGAAGTGG
CACTTGTTATCTGAGCTTGAGGCCGTCCCTACCTGCCCCAGGAGGAGAAGTCTCCATTGTTTCTGTACAACGT
GAAGGGCTACCTGAAGATGGCACCCTCTACCGAATAAACAGATTTAGCTCGATCACAGCACATCCAGAGCGCTGG
GATGTGTCTTCTTCACGGGGGGACCGCTCTGGGCTCTGGACTGGTGCCAGTGCCAGAGGGGGCAGGAGCCTCG
CAATATGTGGCTCTTTTCTCCAGCCCTGACATGAATGAGACACACCCACTGAGCCAGCTTCATTGCGGTCTCTGGG
CTGCTCCAGCTCTGGGGCCTTGGGACCTTGCAAGCAAGAAAGCTGTCTGGCAACAGGGCCACTTTGTCTATGGG
ATTGCTTGTGACAACGGCTGCATCTGGGACCTCAAGTCTGCCCCAGTGGAGCATGGGAACCTCCAGGCACCCCT
CGGAAGGCTCCTCTCCTGCCCCGGTTGGGTCTCTTGGCTCTGCGCTGCTCAGACGGGAAAGTACTGCTATTCACT
CTACCCCATCCGGAGGCCCTGCTGGCTCAGCAACCCCAAGTGCAGTGAAGCCTGCCATATATAAGGTACAATGT
GTGGCAACTCTGCAGGTGGGGTCTATGCAAGCTACAGACCCCTCTGAGTGTGGTCACTGCCTTAGCCTGGCCTGG
ATGCCTACCAGGCCCCACCAACCTAGCTGCTGGATATATAATGGCATGGTGGTTTTCTGGAACCTTCCCACT
AACTACCCCTGCAGCGGATACGGCTCTCTGATGGCTCCTTAAAGCTCTACCCCTTCCAGTGTTCCTAGCCCAT
GACCAGGCTGTGCGTACCCCTTCAATGGTGCAAAGCTAACAGCCATTTCTTGTCTCTGCGGGGAGTGACCGGAAA
ATCAAATTCTGGGACCTTCGACGTCTTACGAACCCATAAACTCTATCAAGCGCTTCTTGAGTACAGAACTGGCC
TGGTGTCTTCCCTACCAATGGTGTCACTGTGGCTCAGGACAACCTGCTATGCCTCTTATGGACTCTGTGGGATTCT
TATATTGACGCTGGTTACCTTGGTTTCAAGGCCTACTTCACTGCTCCTCGAAAAGGCACCGTTTGGAGTCTTTCA
GGATCCGACTGGCTTGGGACAATAGCTGCAGGAGATATATCCGGGGAGCTCATTGCTGCTATATTACCAGATATG
GCATGAATCCAATAAATGTCAAGCGACCTGTAGAGCGAAGATTTCTATATATAAAGCAGATCTGATACCGTAT
CAGGACAGTCTGAAAGTCCAGACCACTTCTTCTGCTTCTATCTGGGGTCCCCAACCCCTCCCAAGGCTCGAACTTAC
ACTGAAACTGTCAACCATCACTACTTGGCTCTTTCAAGACACAGATTTGGGTTTCACTTCCATGATCTGCTCCGTAGA
GAACCAATGCTGCGCATGCAGGAGGGAGAGGGGCATTCTCAACTCTGCCTGGACAGGCTGCAGCTGGAGGCTATT
CATAAGGTACGTTTCAGCCCAAACCTGGACTCCTATGGATGGCTGGTATCTGGGGGGCAGTCAGGCTGGTTTGA
ATCCATTTTGTCCGTGGACTCGCCTCCCCACTGGGCCACCGTATGCAGCTTGAAAGCCGAGCCCACTTCAATGCT
ATGTTCCAACCATCCTCCCCCACTAGACGGCTGGCTTCTCTCCAACAGCCATCGCCTTCTGCCCCACTCCCTAG

CCTTGGCCCCACACCAGATCCTTGGAGTGAAGTCGGTCAAGAACAAATGGCCCCCTATGCACAGAGCCATAGGAAC
GGGGGCCCTTCCCTGGACAGTGATCATGCCAGGCTGGACCTTTAGGCCTGCCTCCCCAGGACTCCTTAGATCCCA
CTCTTTCTACAGACTTCTGTGATCACAGCCCCCTGCGGGCAGGGGGGCTCTCCCTCCACCAACTCTCAAGGCTCC
TCAGCCTAAGACTATGGCTCATGAGAAACACTCAGGCCTGACCTAGGCTTGGGAGTCAAACCTGCTCATATTGAGC
ATATTGTTAAGTGGGTAAAGCCAAGTAAAGGTACTGGGTGTTTTTGTGACCACTTGTGAATGGGTGTATGGAGAA
CTGAAAAGGGTATCTGCATGAAGGCTCCTGTCTGACTATTCCAGGATCCAATATTACTGCCTTCTGAACTTCTT
CTTTAGGGTAACCATCATGTATGCCACGAGGGTGATAGTAATTCGTGAGACTGAAGTTGCTTAGAGTACTTCTT
TGACCAAGGAATACCACAGACACCCCTACCGATAGAACAGTGGCTCAGATCTTACTTGCTCCTGCTTACGAAGTAT
TCCCAATCACTGGTCTATCTGACCCTACTTGAACACTCCTGAACAGTCATGTTTTTAAATCTTCTTTATATCA
AGTCAGAGAGTATACTTCTATAAATTTCACTCATGGATGTTAGGAAATCTAGTCATCTTCCCTGTGATTGCCCTG
TTAAGTATTTAACCATAGCTATCATGTGTTTCCCAAATCTTCTTAGATTAAATATCTTCAGTTACTTC

738/6881

FIGURE 689

MDTCGVGYVALGEAGPVGNMTVVDSFGQEVNLQLDVKTSSSEMTSAEASVEMSLPTPLPGFEDSPDQRRLPPEQES
LSRLEQPDLSSEMSKVSKPRASKPGRKRGGTRKGPKRPPQPNPPSAPLVPGLLDQSNPLSTPMPKKRGRKSKAE
LLLLKLSKDLDRPESQSPKRPPEDFETPSGERPRRRAAQVALLYLQELAEELSTALPAPVSCPEGPKVSSPTKPK
KIRQPAACPGGEEVDGAPRDEDFFLQVEAEDVEESEGPSESSEPEPVVPRSTPRGSTSGKQKPHCRGMAPNGLP
NHIMAPVWKCLHLTKDFREQKHSYWEFAEWIPLAWKWHLLSELEAAPYLPQEEKSPLFSVQREGLPEDGTLYRIN
RFSSITAHPERWDVSFFTGGPLWALDWCPVPEGAGASQYVALFSSPDMNETHPLSQLHSGPGLLQLWGLGTLQOE
SCPGNRAHFVYGIACDNGCIWDLKFCPSGAWELPGTPRKAPLLPRLGLLALACSDGKVLLFSLPHPEALLAQOPP
DAVKPAIYKVQCVALTLQVGSMAQATDPSECGQCLSLAWMPTRPHQHLAAGYYNGMVVFWNLPTNSPLQRIRLSDGS
LKLYPFQCFLAHDQAVRTLQWCKANSHFLVSAGSDRKIKFWDLRRPYEPINSIKRFLSTELAWLLPYNGVTVAQD
NCYASYGLCGIHYIDAGYLGFKAYFTAPRKGTVWSLSGSDWLGTIAAGDISGELIAAILPDMALNPINVKRPVER
RFPIYKADLIPYQDSPEGPDHSSASSGVPNPPKARTYTETVNHHYLLFQDIDLGSFHDLLRREPMLRMQEGEGHS
QLCLDRLQLEAIHKVRFSPNLD SYGWLVS GGQSGLVRIHFVRGLASPLGHRMQLESRAHFNAMFQPSSTTRPFG
SPTSHRLLPTP

739/6881
FIGURE 690

GAGGGACGGTGGGGGTGACCCCTCTCCTCTTGGCTTGACAGGAAGCATGGCACTCTGGCGGGCATAACCAGCGGG
CCCTGGCCGCTCACCCGTGGAAGTACAGGTCCTGACAGCTGGGTCCCTGATGGGCCTGGGTGACATTATCTCAC
AGCAGCTGGTGGAGAGGCGGGGTCTGCAGGAACACCAGAGAGGCCGGACTCTGACCATGGTGTCCCTGGGCTGTG
GCTTTGTGGGCCCTGTGGTAGGAGGCTGGTACAAGGTTTTGGATCGGTTTCATCCCTGGCACCACCAAAGTGGATG
CACTGAAGAAGATGTTGTTGGATCAGGGGGGCTTTGCCCCGTGTTTTCTAGGCTGCTTTCTCCCACTGGTAGGGG
CACTTAATGGACTGTCAGCCCAGGACAACCTGGGCCAAACTACAGCGGGATTATCCTGATGCCCTTATCACCAACT
ACTATCTATGGCCTGCTGTGCAGTTAGCCAACCTTCTACCTGGTCCCCCTTCATTACAGGTTGGCCGTTGTCCAAT
GTGTTGCTGTTATCTGGAACCTCTACCTGTCCTGGAAGGCACATCGGCTCTAAGCCTGCCTCACTCCATCGTTTC
CACCTTGCACTGATGCAGCTTGACCCCTGGAACGGTCAGACAACCTCCTCAAAGTGGGCATACCAGTTTCCACGGG
GTTGGGTTGCCGGTCAGAGCTTAAGAGGACTAGCACCTGCAATGCCCCCTCTTCACTCTAAAATGTACACTGACT
GCTTTAGAGCCCTTGATAATAGTCTTATTCCCACCACATACTAGGCACTCCATAAATATCTGTTGAACCTTCATG
ACCTTATCAACTTTACACCCATATCCCAGCAAATGCCACTCATCCCCACTCTTCATAGACACATTTGTTACTCTA
ACCCTGCCTAGGCTTCTGTAGCTCCAGCTCTTTAGAGACTCCCGGAACCCTTTATATGGTGCCTCAGTAAATAT
GTTATTAAATATGTAATCCGG

740/6881

FIGURE 691

MALWRAYQRALAAHPWKVQVLTAGSLMGLGDIISQQLVERRGLQEHQRGRTLTMVSLGCGFVGPVVGWYKVLDR
FIPGTTKVDALKKMLLDQGGFAPCFLGCFPLPLVGALNGLSAQDNWAKLQRDYPDALITNYYLWPAVQLANFYLP
LHYRLAVVQCVAVIWN SYLSWKAHRL

741/6881
FIGURE 692A

CGCCCCCGCCTCTGAGCTCCCTTCCCATTGGCGGCCCTAGTGTTGGAGGACGGGTCCGTCTGCGGGGCCAGCCCT
TTGGGGCCGCGGTGTGCGACTGCCGGGGAAGTGGTGTTCAAACCGGCATGGTCGGCTACCCCGAGGCCCTCACTG
ATCCCTCCTACAAGGCACAGATCTTAGTGCTCACCTATCCTCTGATCGGCAACTATGGCATCCCCCAGATGAAA
TGGATGAGTTCGGTCTCTGCAAGTGGTTTGAATCCTCGGGCATCCACGTAGCAGCACTGGTAGTGGGAGAGTGCT
GTCCTACTCCCAGCCACTGGAGTGCCACCCGCACCCTGCATGAGTGGCTGCAGCAGCATGGCATCCCTGGCTTGC
AAGGAGTAGACACTCGGGAGCTGACCAAGAAGTTGCGGGAACAGGGGTCTCTGCTGGGGAAGCTGGTCCAGAATG
GAACAGAACCTTCATCCCTGCCATTCTTGGACCCCAATGCCCGCCCCCTGGTACCAGAGGTCTCCATTAAAGACTC
CACGGGTATTCAATACAGGGGGTGCCCTCGGATCCTTGCTTTGGACTGTGGCCTCAAGTATAATCAGATCCGAT
GCCTCTGCCAGCGTGGGGCTGAGGTCACTGTGGTACCCTGGGACCATGCACTAGACAGCCAAGAGTATGAGGGTC
TCTTCTTAAGTAATGGGCGTGGTGACCCTGCCTCCTATCCCAGTGTGCTATCCACACTGAGCCGTGTTTTATCTG
AGCCTAATCCCGACCTGTCTTTGGGATCTGCCTGGGACACCAGCTATTGGCCTTAGCCATTGGGGCCAGACTT
ACAAGATGAGATATGGGAACCGAGGCCATAACCAGCCCTGCTTGTGGTGGGCTCTGGGCGCTGCTTTCTGACAT
CCCAGAACCATGGGTTTGTCTGTGGAGACAGACTCACTGCCAGCAGACTGGGCTCCTCTCTTACCAACGCCAATG
ATGGTTCCAATGAAGGCATTGTGCACAACAGCTTGCCCTTTCTTCAGTGTCCAGTTTCACCCAGAGCACCAGCTG
GCCCTTCAGATATGGAACCTGCTTTTCGATATCTTTCTGGAACTGTGAAAGAGGCCACAGCTGGGAACCTGGGG
GCCAGACAGTTAGAGAGCGGCTGACTGAGCGCCTCTGTCCCCCTGGGATTCCCACTCCCGGCTCTGGACTTCCAC
CACCACGAAAGTTCTGATCCTGGGCTCAGGGGGCCTCTCCATTGGCCAAGCTGGAGAATTGACTACTCGGGCT
CTCAGGCAATTAAGGCCCTGAAGGAGGAAAACATCCAGACGTTGCTGATCAACCCCAATATTGCCACAGTGACAG
CCTCCCAGGGGCTGGCCGACAAGTCTATTTTCTTCCATAACACCTCATTATGTAACCCAGGTGATACGTAATG
AACGCCCCGATGGTGTGTTACTGACTTTTGGGGGCCAGACTGCTCTGAAGTGTGGTGTGGAGCTGACCAAGGCCG
GGGTGCTGGCTCGGTATGGGGTCCGGGTCTGGGCACACCAGTGGAGACCATTGAGCTGACCGAGGATCGACGGG
CCTTTGCTGCCAGAATGGCAGAGATCGGAGAGCATGTGGCCCCGAGCGAGGCAGCAAATCTCTTGAACAGGCC
AGGCAGCCGCTGAACGGCTGGGGTACCCTGTGCTAGTGCGTGCAGCCTTTGCCCTGGGTGGCTGGGCTCTGGCT
TTGCCTCTAACAGGGAGGAGCTCTCTGCTCTCGTGGCCCCAGCTTTTGCCCATACCAGCCAAGTGCTAGTAGACA
AGTCTCTGAAGGGATGGAAGGAGATTGAGTACGAGGTGGTGAGAGACGCCATAGGCAACTGTGTACGGTGTGTA
ACATGGAGAATTTGACCCACTGGGCATCCACACTGGTGAGTCCATAGTGGTGGCCCCCTAGCCAGACACTGAATG
ACAGGGAGTATCAGCTCCTGAGGCAGACAGCTATCAAGGTGACCCAGCACCTGGGAATTGTTGGGGAGTGAATG
TGCAGTATGCCTTGAACCTGAGTCTGAGCAGTATTACATCATTGAAGTGAATGCCAGGCTCTCTCGCAGCTCTG
CCCTGGCCAGTAAGGCCACAGGTTATCCACTGGCTTATGTGGCAGCCAAGCTAGCATTGGGCAATCCCTTTGCCTG
AGCTCAGGAATCTGTGACAGGGGTACAGCAGCCTTTGAACCCAGCGTGGATTATTGTGTGGTGAAGATTCCCTC
GATGGGACCTTAGCAAGTTCTTGCAGTACAGCACAAGATTGGGAGCTGCATGAAGAGCGTTGGTGAAGTCATGG
GCATTGGGCGTTCAATTGAGGAGGCCCTCCAGAAGGCCCTGCGCATGGTGGATGAGAAGTGTGTGGGCTTTGATC
ACACAGTGAACCCAGTCAGCGATATGGAGTTGGAGACTCCAACAGATAAGCGGATTTTTGTGGTGGCAGCTGCTT
TGTGGGCTGGTTATTCAGTGGACCGCTGTATGAGCTCACACGCATCGACCGCTGGTTCCCTGCACCGAATGAAGC
GTATCATCGCACATGCCAGCTGCTAGAACACACCGTGGACAGCCTTTGCCGCCAGACCTGCTGCAACAGGCCA
AGGAAGTGGGGATCTGTCCAGCAGTGAACAGATTGACACAGTTGACAGCTGAGTGGCCAGCCCAGACAAATTACC
TATACCTAACGTATTGGGGCACCACCCATGACCTCACCTTTGGAACACCTCATGTCTAGTCTTGGCTCTGGCG
TCTACCGTATTGGCTTAGCGTTGAATTTGACTGGTGTGCTGTAGGCTGCATCCAGCAGCTCCGAAAGATGGGAT
ATAAGACCATTAGTGAATATAACCCAGAGACAGTCAGCACCAGCTATGACATGTGTGATCGACTCTACTTTG
ATGAGATCTCTTTTGGTGGTGATGGACATCTATGAGCTCGAGAACCCTGAAGGTGTGATCTATCCATGGGTG
GACAGCTGCCCAACAACATGGCCATGGCGTTGCATCGGCAGCAGTGCCGGGTGCTGGGCACCTCCCTGAAGCCA
TTGACTCGGCTGAGAACCCTTTCAAGTTTTCCCGGCTCCTTGACACCATTGGTATCAGCCAGCCTCAGTGGAGGG
AGCTCAGTGACCTGAGTCTGCTCGCCAAATCTGCCAGACCGTGGGGTACCCCTGTGTGGTGCAGCCCTCCTATG
TGCTGAGCGGTGCTGCTATGAATGTGGCTTACACGGATGGAGACCTGGAGCGCTTCTGAGCAGCGCAGCAGCCG
TCTCAAAGAGCATCCCGTGGTCATCTCCAAGTTCAATCCAGGAGGCTAAGGAGATTGACGTGGATGCCGTGGCCT
CTGATGGTGTGGTGGCAGCCATCGCCATCTCTGAGCATGTGGAGAATGCAGGTGTGCATTAGGTGATGCGACGC
TGGTGACCCCCCACAGATATCACTGCCAAAACCTGGAGCGGATCAAAGCCATTGTGCATGCTGTGGGCCAGG

742/6881
FIGURE 692B

AGCTACAGGTCACAGGACCCCTTCAATCTGCAGCTCATTGCCAAGGATGACCAGCTGAAAGTTATTGAATGCAACG
TACGTGTCTCTCGCTCCTTCCCCTTCGTTTCCAAGACACTGGGTGTGGACCTAGTAGCCTTGCCACGCGGGTCA
TCATGGGGGAAGAAGTGAACCTGTGGGGCTAATGACTGGTTCTGGAGTCGTGGGAGTAAAGGTGCCTCAGTTCT
CCTTCTCCCGCTTGGCGGGTGTGACGTGGTGTGGGTGTGAAATGACCAGTACTGGGGAGGTGGCCGGCTTTG
GGGAGAGCCGCTGTGAGGCATACCTCAAGGCCATGCTAAGCACTGGCTTTAAGATCCCCAAGAAGAATATCCTGC
TGACCATTGGCAGCTATAAGAACAAAAGCGAGCTGCTCCCACTGTGCGGCTACTGGAGAGCCTGGGCTACAGCC
TCTATGCCAGTCTCGGCACAGCTGACTTCTACACTGAGCATGGCGTCAAGGTAACAGCTGTGGACTGGCACTTTG
AGGAGGCTGTGGATGGTGAGTGCCACCACAGCGGAGCATCCTGGAGCAGCTAGCTGAGAAAACTTTGAGCTGG
TGATTAACCTGTCAATGCGTGGAGCTGGGGGCCGGCTCTCTTTCCTTTGTACCAAGGGCTACCGCACCCGAC
GCTTGGCCGCTGACTTCTCCGTGCCCTAATCATCGATATCAAGTGCACCAACTCTTTGTGGAGGCCCTAGGCC
AGATCGGGCCAGCCCCCTCTTTGAAGGTGCATGTTGACTGTATGACCTCCCAAAGCTTGTGCGACTGCCGGGAT
TGATTGATGTCCATGTGCACCTGCGGGAACAGGTGGGACACATAAGGAGGACTTTGCTTCAGGCACAGCCGCTG
CCCTGGCTGGGGTATCACCATGGTGTGTGCCATGCCTAATACCCGGCCCCCATCATTGACGCCCTGCTCTGG
CCCTGGCCCAAGCTGGCAGAGGCTGGCGCCGGTGCAGCTTTGCGCTATTCTTGGGGCCTCGTCTGAAAATG
CAGGAACCTTGGGCACCGTGGCCGGGTCTGCAGCCGGGCTGAAGCTTTACCTCAATGAGACCTTCTCTGAGCTGC
GGCTGGACAGCGTGGTCCAGTGGATGGAGCATTTGAGACATGGCCCTCCACCTCCCCATTGTGGCTCACGCAG
AGCAGCAAACCGTGGCTGCTGTCTCATGGTGGCTCAGCTCACTCAGCGCTCAGTGCACATATGTACGTGGCAC
GGAAGGAGGAGATCCTGCTAATTAAGCTGCAAAGGCAGGGGCTTGCCAGTGACCTGCGAGGTGGCTCCCCACC
ACCTGTTCTTAAGCCATGATGACCTGGAGCGCTGGGGCCTGGGAAGGGGAGGTCCGGCCTGAGCTTGGCTCCC
GCCAGGATGTGGAAGCCCTGTGGGAGAACATGGCTGTCTCATGCTTTGCTTGCCTCAGACCATGCTCCCCATACCT
TGGAGGAGAAGTGTGGGTCCAGGCCCCACCTGGGTTCAGAGGTTAGAGACCATGCTGCCACTACTCCTGACGG
CTGTAAGCGAGGGCCGGCTCAGCCTGGACGACCTGCTGCAGCGATTGCACCACAATCCTCGGCGCATCTTTCACC
TGCCCCCGCAGGAGGACACCTATGTGGAGGTGGATCTGGAGCATGAGTGGACAATTCAGCCACATGCCCTTCT
CCAAGGCCCACTGGACACCTTTTGAAGGGCAGAAGTGAAGGGCACCGTCCGCCGTGTGGTCTGCGAGGGGAGG
TTGCCTATATCGATGGGCAGGTCTGTTACCCCGGGCTATGGACAGGATGTACGGAAGTGGCCACAGGGGGCTG
TTCTCAGCTCCACCCCTCAGCCCTGCCACTAGTGAGATGACCACGACACCTGAAAGACCCCGCGTGGCATCC
CAGGGCTTCTGATGGCCGCTTCCATCTGCGGCCCGAATCCATCGAGCCTCCGACCCAGGTTTGCCAGCTGAGG
AGCCAAAGGAGAAGTCTCTCGGAAGGTAGCCGAGCCAGAGCTGATGGGAACCCCTGATGGCACCTGCTACCCTC
CACCACAGTACCAGACAGGCATCTCCCCAGAACCTGGGGACCCCTGGCTTGTGACCCCCAGACCTCACCCC
TGCTGCACTCATTAGTGGGCCAACAATATCCTGTCCGTCCAGCAGTTCACCAAGGATCAGATGTCTCACCTGTTCA
ATGTGGCACACACACTGCGTATGATGGTGCAGAAGGAGCGGAGCCTCGACATCCTGAAGGGGAAGGTATGGCCT
CCATGTTCTATGAAGTGAGCACACGACCAGCAGCTCCTTTGCAGCAGCCATGGCCCGGTGGGAGGTGCTGTGC
TCAGCTTCTCGGAAGCCACATCGTCCGTCCAGAAGGGCGAATCCCTGGCTGACTCCGTGCAGACCATGAGCTGCT
ATGCCGACGTGCTGCTCCGGCACCCCGAGCCTGGAGCAGTGGAGCTGGCCGCCAAGCACTGCCGGAGGCCAG
TGATCAATGCTGGGGATGGGGTCGGAGAGCACCCACCCAGGCCCTGCTGGACATCTTCACCATCCGTGAGGAGC
TGGGAACCTGTCAATGGCATGACGATCAGCATGGTGGGTGACCTGAAGCACGGACGCACAGTACATTCCCTGGCCT
GCCTGCTCACCCAGTATCGTGTGACCTGCGCTACGTGGCACCTCCAGCCTGCGCATGCCACCCACTGTGCGGG
CCTTCGTGGCCTCCCGCGGCACCAAGCAGGAGGAATTCGAGAGCATTGAGGAGGCGCTGCCTGACACTGATGTGC
TCTACATGACTCGAATCCAGAAGGAACGATTTGGCTCTACCCAGGAGTACGAAGCTTGCTTTGCTCAGTTTCATCC
TCACTCCCCACATCATGACCCGGGCCAAGAAGAAGATGGTGGTGTGACCCGATGCCCCGTGTCAACGAGATAA
GCGTGAAGTGGACTCGGATCCCCGCGCAGCCTACTTCCGCCAGGCTGAGAACGGCATGTACATCCGCATGGCTC
TGTTAGCCACCGTGTGGGCCGTTTCTAGGGGCTGGCTTCTCAGCCTCTTCTCTTTAGGCCAGCTGCTGGGC
AAGGAATTCCAGTGCCTCCTACGGGGGCAGCACACTTAGATATTCCTGGACATCCAGATTGCTCAGATGTGCTGA
CCACACTTCAGGCTCTGGACTGGAGCTCTCTGGCATGGGGTGGGGCCTCAGATGCTGGGGCCAGCTGCCCCA
TCTTCATTCTGCACCTTAAACCTGTACAGTCATTTTCTACTGACTTAATAAACAGCCGAGCTGTCCCTTG

743/6881
FIGURE 693

MAALVLEDGSVLRGQPFGAAVSTAGEVVFQTMVGYPEALTDPSYKAQILVLTYPPLIGNYGIPPEMDEFGLCKW
FESSGIHVAALVVGECPTPSHWSATRTLHEWLQQHGIPGLQGVDTRELTKKLREQGSLLGKLVQNGTEPSSLPF
LDPNARPLVPEVSIKIPRVFNTGGAPRI LALDCGLKYNQIRCLCQRGAETVVPWDHALDSQEYEGFLFSLNGPGD
PASYPVSVVSTLSRVLSEPNRPVFGICLGHQLLALAIGAKTYKMRYGNRGHNQPCLLVGSGRCFLTSONHGFAVE
TDSL PADWAPLFTNANDGSNEGIVHNSLPFFSVQFHPEHQAGPSDMELLFDIFLETVKEATAGNPGGQTVRERLT
ERLCPPGIPTPGSGLPPPRKVLILGSGGLSIGQAGEFDYSGSQAICALKEENIQTLINPNIA TVQTSQGLADKV
YFLPITPHYVTQVIRNERPDGVLLTFGGQTALNCGVELTKAGVLARYGVRVLGTPVETIELTEDRRFAAARMAEI
GEHVAPSEAANSLEQAQAAERLGPVLRVRAAFALGGLSGGFASNREELSALVAPAFAHTSQVLVDKSLKGWKEI
EYEVVRDAYGNCVTVCNMENLDPLGIHTGESIVVAPSQTINDREYQLLRQTAIKVTQHLGIVGECNVQYALNPES
EQYYIIIEVNARLSRSSALASKATGYPLAYVAAKLALGIPPELRNSVTGGTAAFEPSVDYCVVKIPRWDLSKFLR
VSTKIGSCMKSVGEVMGIGRSFEEAFQKALRMVDENCVGFDHTVKPVSDMELETPTDKRIFVVAALWAGYSVDR
LYELTRIDRWFLHRMKRIIAHAQILLEQHRGQPLPPDLLQQAACLGFSQKIALAVLSTELAVRKLQELGICPAV
KQIDTVAAEWPAQTNYLYLTYWGTTHDLTFRTPHVLVLSGSGVYRIGSSVEFDWCAVGCIQQLRKMGYKTIMVNYN
PETVSTDYDMCDRLYFDEISFEVMDIYELENPEGVILSMGGQLPNNMAMALHRQQCRVLGTSPEAIDSAENRFK
FSRLLDITIGISQPPQWRELSDESARQFCQTVGYPCVVRPSYVLSGAAMNVAYTDGDLERFLSSAAAVSKEHPVVI
SKFIEAKEIDVDVAVASDGVVAAIAISEHVENAGVHSGDATLVTPPDITAKTLERIKAIHVAVGQELQVTGPFN
LQLIAKDDQLKVIENVRVRSRFPFVSKTLGVDLVALATRVIMGEEVEPVGLMTGSGVVGKVPQFSF SRLAGAD
VVLGVEMTSTGEVAGFGESRCEAYLKAMLSTGFKIPKKNILLTIGSYKNKSELLPTVRLLESLSGLYSLYASLGAD
FYTEHGVKVTAVDWHFEEAVDGECPQRSILEQLAEKNFELVINLSMRGAGGRRLSSSVTKGYRTRRLAADFSVP
LIIDIKCTKLFVEALGQIGPAPPLKVHVD CMTS QKLVRPLGLIDVHVHLREP GGTHKEDFASGTAAALAGGITMV
CAMPNTRPPIIDAPALALAQKLAEGARCDFALFLGASSENAGTLGT VAGSAAGLKL YLNETFSELRLDSVVQWM
EHFETWPSHLP IVAHAEQQTVA AVLMAQLTQRSVHICHVARKEEILLIKAARGLPVTCEVAPHHLFLSHDDL
ERLGP GKGEVRPELGS RDVEALWENMAVIDCFASDHAPHTLEEKCSRPPPGFPGLTMLPLLLTAVSEGRSL
DDLQRLHHPNRIFHLPPQEDTYVEVDLEHEWTIPSHMPFSKAHWTPFEGQKVKGTVRRVLRGEVAYIDQVL
VPPGYGQDVRKWPQGA VPQLPPSAPATSEMTTTPRPRRGIPGLPDGRFHLPPRIHRASDPGLPAEEPKEKSSRK
VAEPELMGTPDGTCTYPPPPVPRQASPNLGTPLHLPQTSPLHSLVGHILSVQQFTKDQMSHLFNVAHTLRMM
VQKERSLDILKGKVMASMFYEVSTRTSSSF AAAMARLGAVLSFSEATSSVQGESLADSVQTMSCYADV VVLRH
PQPGAVELAAKHCRRPVINAGDGVGEHPTQALLDIFTIREELGTVNGMTITMVGDLKHGRTVHSLACLLTQYRVS
LRYVAPPSLRMPPTVRAFVASRGTKQEEFESIEEALPD TDVLYMTRIQKERFGSTQEYEACFGQFILT PHIMTRA
KKKMVMHMPRVNEISVEVDS DPRAAYFRQAENGYIR MALLATVLRG

744/6881
FIGURE 694

GGAGCACCAAGGGAACGGAAAATGGCGCCTCACGGCCCGGGTAGTCTTACGACCCTGGTGCCCTGGGCTGCCGCC
CTGCTCCTCGCTCTGGGCGTGGAAGGGCTCTGGCGCTACCCGAGGTACAGAAGCAAGTTTGAGGTCGGGCTGAA
GCAGGGTCACTGGCCAGCCGTGCGTCGCGCTCGCCAGCGGCTCCCCCTTCTCCTCGGCGGGCCTGCGGTTCTGAT
TTCGTCCCTGACGCTTCCCGACCCTGCCCAGCCAGATATGCACCCAATGTCCAGGGAGCGTGCAAAATTGTCAA
AAGTGGCCTTTTATTGTAAAACGACACGAGAGCTAATGCTGCATGCCCGTTGCTGCCTGAATCAGAAGGGCACCA
TCTTGGGGCTGGATCTCCAGAACTGTTCTCTGGAGGACCCTGGTCCAACTTTTCATCAGGCACATACCACTGTCA
TCATAGACCTGCAAGCAAACCCCTCAAAGGTGACTTGGCCAACACCTTCCGTGGCTTTACTCAGCTCCAGACTC
TGATACTGCCACAACATGTCAACTGTCCTGGAGGAATTAATGCCTGGAATACTATCACCTCTTATATAGACAACC
AAATCTGTCAAGGGCAAAAGAACCTTTGCAATAACACTGGGGACCCAGAAATGTGTCCTGAGAATGGATCTGTG
TACCTGATGGTCCAGGTCTTTTGCAGTGTGTTTGTGCTGATGGTTTCCATGGATACAAGTGTATGCGCCAGGGCT
CGTTCTCACTGCTTATGTTCTTCGGGATTCTGGGAGCCACCACTCTATCCGTCTCCATTCTGCTTTGGGCGACCC
AGCGCCGAAAAGCCAAGACTTCATGAACACTACATAGGTCTTACCATTGACCTAAGATCAATCTGAACTATCTTAGC
CCAGTCAGGGAGCTCTGCTTCCTAGAAAGGCATCTTTCGCCAGTGGATTGCGCTCAAGGTTGAGGCCGCCATTGG
AAGATGAAAATTGCACTCCCTTGGTGTAGACAAATACCAGTTCCTTGGTGTGTTGCCTATAATAAACACTT
TTTCTTTTTAAAAA

745/6881

FIGURE 695

MLHARCCLNQKG TILGLDLQNC SLEDPGPNF HQAHTTVIIDLQANPLK GDLANTFRGFTQLQTLILPQHVNCPGG
INAWNTITSYIDNQICQGQKNLCNNTGDPEMCPENGSCVPDGPGLLCVCADGFHGYKCMRQGSFSLMFFGILG
ATTLSVSILLWATQRRKAKTS

746/6881
FIGURE 696

GGAGCACCAAGGGAACGGAAAAATGGCGCCTCACGGCCCCGGGTAGTCTTACGACCCTGGTGCCCTGGGCTGCCGCC
CTGCTCCTCGCTCTGGGCGTGGAAGGGCTCTGGCGCTACCCGAGATATGCACCCAATGTCCAGGGAGCGTGCAA
AATTTGTCAAAAGTGGCCTTTTATTGTAAAACGACACGAGAGCTAATGCTGCATGCCCGTTGCTGCCTGAATCAG
AAGGGCACCATCTTGGGGCTGGATCTCCAGAACTGTTCTCTGGAGGACCCTGGTCCAACTTTTCATCAGGCACAT
ACCACTGTCATCATAGACCTGCAAGCAAACCCCTCAAAGGTGACTTGGCCAACACCTTCCGTGGCTTTACTCAG
CTCCAGACTCTGATACTGCCACAACATGTCAACTGTCCTGGAGGAATTAATGCCTGGAATACTATCACCTCTTAT
ATAGACAACCAAATCTGTCAAGGGCAAAGAACCTTTGCAATAACACTGGGGACCCAGAAATGTGTCCTGAGAAT
GGATCTTGTGTACCTGATGGTCCAGGTCTTTTGCAGTGTGTTTGTGCTGATGGTTTCCATGGATACAAGTGTATG
CGCCAGGGCTCGTTCTCACTGCTTATGTTCTTCGGGATTCTGGGAGCCACCACTCTATCCGTCTCCATTCTGCTT
TGGGCGACCCAGCGCCGAAAAGCCAAGACTTCATGAACACTACATAGGTCTTACCATTGACCTAAGATCAATCTGAA
CTATCTTAGCCCAGTCAGGGAGCTCTGCTTCCTAGAAAGGCATCTTTCGCCAGTGGATTCCCTCAAGGTTGAGG
CCGCCATTGGAAGATGAAAAATTGCACTCCCTTGGTGTAGACAAATACCAGTTCCTTGGTGTGTGCTGCTATA
ATAACACTTTTTCTTTTAAAAA

747/6881
FIGURE 697

MLHARCCLNQKGTILGLDLQNCSEDPGPNFHAHTTVIIDLQANPLKGDLANFRGFTQLQTLILPQHVNCPGG
INAWNTITSYIDNQICQGQKNLCNNTGDPPEMCPENGSCVPDGPGLLQCVCADGFHGYKCMRQGSFSLMFFGILG
ATTLSVSILLWATQRRKAKTS

748/6881
FIGURE 698

CGGCCCTAGCCCTCTTTTCGGGGATACTGGCCGACCCCCCTCTTCCCTTTTCCCTTTAGTGAAGGCCTCCCCCGTCG
CCGCGCGGGCTTCCCGGAGCCGACTGCAGACTCCCTCAGCCCGGTGTTCCCGCGTCCGGACGCCGAGGTCGCGGC
TTCGCAGAACTCGGGCCCCCTCCATCCGCCCTCAGAAAAGGGAGCGATGTTGATCTCAGGAAGCACAAAGGGACC
TTCCTAGCTCTGACTGAACCACGGAGCTCACCTGGACAGTATCACTCCGTGGAGGAAGACTGTGAGACTGTGGC
TGAAGCCAGATTGTAGCCACACATCCGCCCTGCCCTACCCAGAGCCCTGGAGCAGCAACTGGCTGCAGATCA
CAGACACAGTGAGGATATGAGTGTAGGGGTGAGCACCTCAGCCCCCTTTTCCCAACCTCGGGCACAAAGCGTG
CATGTCTACCTTCTCCATCATGGACTATGTGGTGTGCTGCTGCTGGTTCCTCTCTTGGCATTGGGCTCTA
CCATGCTTGTGCTGGCTGGGGCCGGCATACTGTTGGTGAGCTGCTGATGGCGGACCGCAAAATGGGCTGCCTTCC
GGTGGCACTGTCCCTGCTGGCCACCTTCCAGTCAGCCGTGGCCATCCTGGGTGTGCCGTGAGATCTACCGATT
TGGGACCCAATATTGGTTCCCTGGGCTGCTGCTACTTTCTGGGGCTGCTGATACCTGCACACATCTTATCCCCGT
TTTCTACCGCTGCATCTCACCAGTGCCTATGAGTACCTGGAGCTTCGATTCAATAAACTGTGCGAGTGTGGG
AACTGTGACCTTCATCTTTTCAGATGGTGATCTACATGGGAGTTGTGCTCTATGCTCCGTCAATTGGCTCTCAATGC
AGTGAAGGCGGTGATCTGTGGCTGTCCGTGCTGGCCCTGGGCATTGTCTGTACCGTCTATACAGCTCTGGGTGG
GCTGAAGGCGGTGATCTGGACAGATGTGTTCCAGACACTGGTTCATGTTCCCTCGGGCAGCTGGCAGTTATCATCGT
GGGGTCAGCCAAGGTGGGCGGCTTGGGGCGTGTGTGGGCGGTGGCTTCCAGCACGGCCGATCTCTGGGTTTGA
GCTGGATCCAGACCCCTTTGTGCGGCACACCTTCTGGACCTTGGCCTTCGGGGGTGTCTTCATGATGCTCTCCTT
ATACGGGGTGAACCAGGCTCAGGTGCAGCGGTACCTCAGTTCGCCGACGGAGAAGGCTGCTGTGCTCTCCTGTTA
TGCAGTGTTCCTTCCAGCAGGTGTCCCTCTGCGTGGGCTGCCTCATTGGCCTGGTTCATGTTTCGCGTATTACCA
GGAGTATCCCATGAGCATTACAGAGGCTCAGGCAGCCCCAGACAGTTCGTCTGTACTTTGTGATGGATCTCCT
GAAGGGCTGCCAGGCTGCCAGGCTCTTCATTGGCTGCCTCTTCAGCGGCTCTCTCAGCACTATATCCTCTGC
TTTTAATTCAATTGGCAACTGTTACGATGGAAGACCTGATTTCGACCTTGGTTCCCTGAGTTCTCTGAAGCCCCGGC
CATCATGCTTTCCAGAGGCTTGCCTTTGGCTATGGGCTGCTTTGTCTAGGAATGGCCTATATTTCTCCAGAT
GGGACCTGTGCTGCAGGCAGCAATCAGCATCTTTGGCATGGTTGGGGACCGCTGCTGGGACTCTTCTGCCTTGG
AATGTTCTTTCCATGTGCTAACCTCCTGGTGTGTTGTGGGCTGTTGGCTGGGCTCGTCATGGCCTTCTGGAT
TGGCATCGGGAGCATCGTGACCAGCATGGGCTTCAGCATGCCACCTCTCCCTCTAATGGGTCCAGCTTCTCCCT
GCCCACCAATCTAACCGTTGCCACTGTGACCACACTGATGCCCTTGACTACCTTCTCCAAGCCACAGGGCTGCA
GCGGTTCTATTCTTGTCTTACTTATGTTACAGTGCTCACAACCTCCACCACAGTGATTGTGGTGGGCTGATTGT
CAGTCTACTCACTGGGAGAATGCGAGGCGGCTCCCTGAACCTGCAACCATTTACCCAGTGTGCCAAAGCTCCT
GTCCCTCCTTCCGTTGCTGCTCAGAAAGCGGCTCCACTGCAGGAGCTACGGCCAGGACCACCTCGACACTGGCCT
GTTTCTGAGAAGCCGAGGAATGGTGTGCTGGGGACAGCAGAGACAAGGAGGCCATGGCCCTGGATGGCACAGC
CTATCAGGGGAGCAGCTCCACCTGCATCCTCCAGGAGACCTCCCTGTGATGTTGACTCAGGACCCCGCTCTGTC
CTCACTGTGCCAGGCCATAGCCAGAGGCCACCCTGTAGTACAGGGATGAGTCTTGGTGTGTTCTGCAGGGACAGG
CCTGGATGATCTAGCTCATACCAAGGACCTTGTCTGAGAGGTTCTTGCCTGCAGGAGAAGCTGTACATCTCA
AGCATGTGAGGCACCGTTTTTCTCGTCGCTTGCCAATCTGTTTTTAAAGGATCAGGCTCGTAGGGAGCAGGATC
ATGCCAGAAATAGGGATGGAAGTGATCCTCTGGGAAAAAGATAATGGCTTCTGATTCAACATAGCCATAGTCCT
TTGAAGTAAGTGGCTAGAAACAGCACTCTGGTTATAATTGCCCCAGGGCTGATTACAGGACTGACTCTCCACCAT
AAAAGTGAAGCTGCTTCCCTGTAGTCCCATTTTTCAGTACCAGTTCTGCCAGCCACAGTGAGCCCTATTATTA
CTTTCAGATTGTCTGTGACACTCAAGCCCTCTCATTTTTATCTGTCTACCTCCATTCTGAAGAGGGAGGTTTTG
GTGTCCCTGGTCTCTGCGAATAGAAGATCCATTTGTCTTTGTGTAGAGCAAGCACGTTTTCCACCTCACTGTCT
CCATCCTCCACCTCTGAGATGGACACTTAAGAGACGGGGCAATGTGGATCCAAGAAACCAGGGCCATGACCAGG
TCCACTGTGGAGCAGCCATCTATCTACCTGACTCCTGAGCCAGGCTGCCGTGGTGTCAATTTCTGTCATCCGTGCT
CTGTTTCTTTTGGAGTTTCTTCTCCACATTATCTTTGTTCTGGGGAATAAAACTACCATTGGACCTAAAAAA
AAAAAAAAAAAA

749/6881

FIGURE 699

MSVGVSTSAPLSPTSGTSVGMSTFSIMDYVVFVLLLVLSLAIGLYHACRGWGRHTVGE LLMADRKMGC L PVALSL
LATFQSAVAILGVPSEIYRFGTQYWFLGCCYFLG LLIPAHIFIPVFYRLHLTSAYEYLELRFNKTVRVC GTVTFI
FQMVIYMGV VLYAPSLALNAV TGF DLWLSVLALGIVCTVYTALGGLKAVIWT DVFQTLVMFLGQLAVIIVGSAKV
GGLGRVWAVASQHGRI SGFELDPDFVRHTFWTLAFGGVFMMLSLYGVNQAQVQRYLSSRTEKAAVLSCYAVFPF
QQVSLCVGCLIGLVMFAYYQEYPMSIQQAQAAPDQFVLYFVMDLLKGLPGLPGLFIACLFSGSLSTISSAFNSLA
TVT MEDLIRPWPEFSEARAIMLSRGLAFGYGLLCLGMAYISSQMGPVLQAAISIFGMVGGPLLGLFCLGMFFPC
ANPPGAVVG LLAGLVMAFWIGIGSIVTSMGFSMPPSPSNGSSFSLPTNLT VATVTTL MPLTTFSKPTGLQR FYSL
SYLWYSAHNSTTVIVVGLIVSLLTGRMRGRSLNPAT IYPVLPKLLSLLPLSCQKRLHCRSYGQDHLD TGLFPEKP
RNGVLGDSRDKEAMALDGTAYQGSSSTCILQETSL

751/6881
FIGURE 701

GTAGCCTCATGGGAAGAGAAGCAGATCCTGTGCGTGGGGCTAGTGGTGCTGGACGTCATCAGCCTGGTGGACAAGT
ACCCTAAGGAGGACTCGGAGATAAGGTGTTTGTCCCAGAGATGGCAGCGCGGAGGCAACGCGTCCAACCTCCTGCA
CCGTTCTCTCCCTGCTCGGAGCCCCCTGTGCCTTCATGGGCTCAATGGCTCCTGGCCATGTTGCTGATTTTGTCC
TGGATGACCTCCGCCGCTATTCTGTGGACCTACGCTACACAGTCTTTCAGACCACAGGCTCCGTCCCCATCGCCA
CGGTCAATCATCAACGAGGCCAGTGGTAGCCGCACCATCCTATACTATGACAGGAGCCTGCCAGATGTGTCTGCTA
CAGACTTTGAGAAGGTTGATCTGACCCAGTTCAAGTGGATCCACATTGAGGGCCGGAACGCATCGGAGCAGGTGA
AGATGCTGCAGCGGATAGACGCACACAACACCAGGCAGCCTCCAGAGCAGAAGATCCGGGTGTCCGTGGAGGTGG
AGAAGCCACGAGAGGAGCTCTTCAGCTGTTTGGCTACGGAGACGTGGTGTTTGTGAGCAAAGATGTGGCCAAGC
ACTTGGGGTTCCAGTCAGCAGAGGAAGCCTTGAGGGGCTTGTATGGTCGTGTGAGGAAAGGGGCTGTGCTTGTCT
GTGCCTGGGCTGAGGAGGGCGCCGACGCCCTGGGCCCTGATGGCAAATTGCTCCACTCGGATGCTTTCCCGCCAC
CCCGCGTGGTGGATACACTGGGAGCTGGAGACACCTTCAATGCCTCCGTCATCTTCAGCCTCTCCAGGGGAGGA
GCGTGCAGGAAGCACTGAGATTCGGGTGCCAGGTGGCCGGCAAGAAGTGTGGCCTGCAGGGCTTTGATGGCATCG
TGTGAGAGCAGGTGCCGGCTCCTCACACACCATGGAGACTACCATTGCGGCTGCATCGCCTTCTCCCTCCATCC
AGCCTGGCGTCCAGGTTGCCCTGTTTCAGGGGACAGATGCAAGCTGTGGGGAGGACTCTGCCTGTGTCTGTGTTT
CCCACAGGGAGAGGCTCTGGGGGGATGGCTGGGGGATGCAGAGCCTCAGAGCAAATAAATCTTCTCAGAGCCAG
CTTCTCCTCTCAATGTCTGAACTGCTCTGGCTGGGCATTCTGAGGCTCTGACTCTTCGATCCTCCCTCTTTGTG
TCCATTCCCCAAATTAACCTCTCCGCCCAGGCCAGAGGAGGGGCTGCCTGGGCTAGAGCAGCGAGAAGTGCCCT
GGGCTTGCCACCAGCTCTGCCCTGGCTGGGGAGGACACTCGGTGCCCCACACCCAGTGAACCTGCCAAAGAAACC
GTGAGAGCTCTTCGGGGCCCTGCGTTGTGCAGACTCTATTCCCACAGCTCAGAAGCTGGGAGTCCACACCGCTGA
GCTGAAGTGCAGAGCCAGTGGGGGGCAGGGGTGCGCCTCCTCTGCCCTGCCACCAGCCTGTGATTTGATGGGGT
CTTCATTGTCCAGAAATACCTCCTCCCGCTGACTGCCCCAGAGCCTGAAAGTCTCACCTTGGAGCCCCACCTTGG
AATTAAGGGCGTGCCTCAGCCACAAATGTGACCCAGGATACAGAGTGTGCTGTCTCAGGGAGGTCCGATCTGG
AACACATATTGGAATTGGGGCCAACTCCAATATAGGGTGGGTAAGGCCTTATAATGTAAAGAGCATATAATGTAA
AGGGCTTTAGAGTGAGACAGACCTGGATTCAAATCTGCCATTTAATTAGCTGCATATCACCTTAGGGTACAGCAC
TTAACGCAATCTGCCTCAATTTCTTCATCTGTCAAATGGAACCAATTCTGCTTGGCTACAGAATTATTGTGAGGA
TAAAAATCATATATAAAAAAAAAA

752/6881
FIGURE 702

MEEKQILCVGLVLDVISLVDPKPKEDSEIRCLSQRWQRGGNASNSCTVLSLLGAPCAFMGSMAPGHVADFVLDD
LRRYSVDLRYTVFQTTGSVPATVIINEASGSRTILYYDRSLPDVSATDFEKVDLTQFKWIHIEGRNASEQVKML
QRIDAHNTRQPPEQKIRVSVEVEKPREELFQLFGYGDVVFVSKDVAKHLGFQSAEEALRGLYGRVRKGAVLVCAW
AEEGADALGPDGKLLHSDAFPPPRVDTLGAGDTFNASVIFSLSQGRSVQEALRFGCQVAGKKCGLQGF DGIV

753/6881
FIGURE 703

GTAGCCTCATGGAAGAGAAGCAGATCCTGTGCGTGGGGCTAGTGGTGCTGGACGTCATCAGCCTGGTGGACAAGT
ACCCTAAGGAGGACTCGGAGATAAGGTGTTTGTCCCAGAGATGGCAGCGCGGAGGCAACGCGTCCAACTCCTGCA
CCGTTCTCTCCCTGCTCGGAGCCCCCTGTGCCTTCATGGGCTCAATGGCTCCTGGCCATGTTGCTGACTTCCTGG
TGGCCGACTTCAGGCGGCGGGGCGTGGACGTGTCTCAGGTGGCCTGGCAGAGCAAGGGGGACACCCCCAGCTCCT
GCTGCATCATCAACAACCTCCAATGGCAACCGTACCATTGTGCTCCATGACACGAGCCTGCCAGATGTGTCTGCTA
CAGACTTTGAGAAGGTTGATCTGACCCAGTTCAAGTGGATCCACATTGAGGGCCGGAACGCATCGGAGCAGGTGA
AGATGCTGCAGCGGATAGACGCACACAACACCAGGCAGCCTCCAGAGCAGAAGATCCGGGTGTCCGTGGAGGTGG
AGAAGCCACGAGAGGAGCTCTTCCAGCTGTTTGGCTACGGAGACGTGGTGTGTTGTCAGCAAAGATGTGGCCAAGC
ACTTGGGGTTCCAGTCAAGCAGAGGAAGCCTTGAGGGGCTGTATGGTTCGTGTGAGGAAAGGGGCTGTGCTTGTCT
GTGCTTGGGCTGAGGAGGGCGCCGACGCCCTGGGCCCCTGATGGCAAATTGCTCCACTCGGATGCTTTCCCGCCAC
CCCGCTGGTGGATACACTGGGAGCTGGAGACACCTTCAATGCCTCCGTCATCTTCAGCCTCTCCAGGGGAGGA
GCGTGCAGGAAGCACTGAGATTCCGGTGCCAGGTGGCCGGCAAGAAGTGTGGCCTGCAGGGCTTTGATGGCATCG
TGTGAGAGCAGGTGCCGGCTCCTCACACACCATGGAGACTACCATTGCGGCTGCATCGCCTTCTCCCCTCCATCC
AGCCTGGCGTCCAGGTTGCCCTGTTTCAAGGGACAGATGCAAGCTGTGGGGAGGACTCTGCCTGTGTCTGTGTTT
CCCACAGGGAGAGGCTCTGGGGGGATGGCTGGGGGATGCAGAGCCTCAGAGCAAATAAATCTTCTCAGAGCCAG
CTTCTCCTCTCAATGCTGAAGTGTCTGGCTGGGCATTCTGAGGCTCTGACTCTTCGATCCTCCCTCTTTGTG
TCCATTCCCCAAATTAACCTCTCCGCCCAGGCCAGAGGAGGGGCTGCCTGGGCTAGAGCAGCGAGAAGTGCCCT
GGGCTTGCCACCAGCTCTGCCCTGGCTGGGGAGGACACTCGGTGCCCCACACCCAGTGAACCTGCCAAAGAAACC
GTGAGAGCTCTTCGGGGCCCTGCGTTGTGCAGACTCTATTCCACAGCTCAGAAGCTGGGAGTCCACACCGCTGA
GCTGAACTGACAGGCCAGTGGGGGGCAGGGGTGCGCCTCCTCTGCCCTGCCACCAGCCTGTGATTGATGGGGT
CTTCATTGTCCAGAAATACCTCCTCCCGCTGACTGCCCCAGAGCCTGAAAGTCTCACCTTGGAGCCACCTTGG
AATTAAGGGCGTGCCTCAGCCACAAATGTGACCCAGGATACAGAGTGTGCTGTCCTCAGGGAGGTCCGATCTGG
AACACATATTGGAATTGGGGCCAACTCCAATATAGGGTGGGTAAGGCCTTATAATGTAAAGAGCATATAATGTAA
AGGGCTTTAGAGTGAGACAGACCTGGATTCAAATCTGCCATTTAATTAGCTGCATATCACCTTAGGGTACAGCAC
TTAACGCAATCTGCCTCAATTCTTCATCTGTCAAATGGAACCAATTCTGCTTGGCTACAGAATTATTGTGAGGA
TAAAAATCATATATAAAAAAAAAA

754/6881
FIGURE 704

MEEKQILCVGLVVLDVISLVDPKEDSEIRCLSQRWQRGGNASNSCTVLSLLGAPCAFMGSMAPGHVADFLVAD
FRRRGVDVSQVAWQSKGDTTPSSCCIINNSNGNRTIVLHDTSLPDVSATDFEKVDLTQFKWIHIEGRNASEQVKML
QRIDAHNTRQPPEQKIRVSVEVEKPREELFQLFGYGDVVFVSKDVAKHLGFQSAEEALRGLYGRVRKGAVLVCAW
AEEGADALGPDGKLLHSDAFPPRVDTLGAGDTFNASVIFSLSQGRSVQEALRFGCQVAGKKCGLQGFDGIV

755/6881
FIGURE 705

ATGGGCCCCCGCACCCCTCTGGAGCTGCTACCTCTGCTGCCTGCTGACGGCAGCTGCAGGGGCCGCCAGCTACCCCT
CCTCGAGGTTTCAGCCTCTACACAGGTTCCAGTGGGGCCCTCAGCCCCGGGGGGCCCCAGGCCAGATTGCCCCC
CGGCCAGCCAGCCGCCACAGGAAGTGGTGTGCCTACGTGGTGACCCGGACAGTGAGCTGTGTCTTGGAGATGGA
GTGGAGACATATGTCAAGTACCAGCCTTGTGCCTGGGGCCAGCCCCAGTGTCCCCAAAGCATCATGTACCGCCGC
TTCCTCCGCCCTCGCTACCGTGTGGCCTACAAGACAGTGACCGACATGGAGTGGAGGTGCTGTACAGGGTTATGGG
GGCGATGACTGTGCTGAGAGTCCCGCTCCAGCGCTGGGGCCTGCGTCTTCCACACCACGGCCCCCTGGCCCCGCT
GCCCCGCCCAACCTCTCTGGCTCCAGTGCAGGCAGCCCCCTCAGTGGACTGGGGGGAGAAGGTCTGGGGAGTCA
GAGAAGGTGCAGCAGCTGGAGGAACAGGTGCAGAGCCTGACCAAGGAGCTGCAAGGCTGCGGGGCGTCTGCAA
GGACTGAGCGGGCGCCTGGCAGAGGATGTGCAGAGGGCTGTGGAGACGGCCTTCAACGGGAGGCAGCAGCCAGCT
GACGCGGCTGCCCCGCCCTGGGGTGCATGAAACCTCAATGAGATCCAGCACCAGCTGCAGCTCCTGGACACCCGC
GTCTCCACCCACGACCAGGAGCTGGGTACCTCAACAACCATCATGGCGGCAGCAGCAGCAGTGGGGGCAGCAGG
GCCCCAGCCCCAGCCTCAGCCCCCTCCGGGGCCCCAGTGAGGAGCTGCTGCGGCAGCTGGAGCAGCGGTTGCAGGAG
TCCTGCTCCGTGTGCTGGCCGGGCTAGATGGCTTCCGCCGGCAGCAGCAGGAGGACAGGGAGCGGCTGCCAGCG
ATGGAGAAGCTGCTGGCCTCGGTGGAGGAGCGGCAACGGCACCTCGCAGGGCTGGCGGTGGGGCCGACGGCCCCCT
CAGGAATGCTGCTCTCCAGAGCTGGGCCGGCGACTGGCAGAGCTGGAGCGCAGGCTGGATGTCTGTGGCCGGCTCA
GTGACAGTGCTGAGTGGGCGGCGAGGCACAGAGCTGGGAGGAGCCGCGGGGCAGGGAGGCCACCCCCAGGCTAC
ACCAGCTTGGCCTCCCGCCTGTCTCGCCTGGAGGACCGCTTCAACTCCACCCTGGGGCCCTTCGGAGGAGCAGGAG
GAGAGCTGGCCTGGGGCTCCTGGGGGGCTGAGCCACTGGCTGCCTGCTGCCCCGGGGCCGACTAGAGCAGTTGGGG
GGGCTGTGGCCAATGTGAGCGGGGAGCTGGGGGGGCGGTTGGATCTGTTGGAGGAGCAGGTGGCAGGGGCCATG
CAGGATGCGGGCAGCTCTGCTCTGGGGCCCCTGGGGAGCAGGACTCTCAAGTCAGCGAGATCCTCAGTGCCTTG
GAGCGCAGGGTGCTGGACAGTGAGGGGCAGCTGCGGCTGGTGGGCTCCGGCCTGCACAGGTGGAAGCAGCGGGG
GAGGCCCGGCAGGCCACGCTGGAGGGATTACAAGAGGTTGTGGGCCGGCTCCAGGATCGTGTGGATGCCCAGGAT
GAGACAGCTGCAGAGTTCACTACGGCTGAATCTCACTGCGGCCCGGCTAGGCCAACTGGAGGGGCTGCTGCAG
GCCCATGGGGATGAGGGCTGTGGGGCCTGTGGCGGAGTCCAAGAGGAAGTGGGCCCGCTTCGGGATGGTGTGGAG
CGCTGCTCCTGCCCCCTGTTGCCCTCCTCGGGTCTGGGGCTGGTCCAGGTGTTGGGGGCCCAAGCCGTGGGGCC
CTGGACGGCTTCAGCGTGTGTTGGGGGCAGCTCAGGCTCAGCCCTGCAGGCCCTGCAAGGAGAGCTCTCTGAGGTT
ATTCTCAGCTTCAGCTCCCTCAATGACTCACTGAATGAGCTCCAGACCACTGTGGAGGGCCAGGGCGCTGATCTG
GCTGACCTGGGGGCAACCAAGGACCGTATCATTCTGAGATTAACAGGCTGCAGCAGGAGGCCACAGAGCATGCT
ACAGAGAGTGAAGAGCGCTTCCGAGGCCTAGAGGAGGACAAGCACAGGCCGGCCAGTGCCCCAGCTTAGAGGGG
CGATTGGGCCGTCTTGAGGGTGTCTGTGAACGGTTGGACACTGTGGCTGGGGGACTGCAGGGCCTGCGCGAGGGC
CTTTCCAGACAGTGGCTGGGCTCTGGGCTGGGCTCCGGGAAACCAACACCACCAGCCAGATGCAGGCAGCCCTG
CTGGAGAAGCTGGTCCGGGGACAGGCCGGGCTGGGCAGGCGGCTGGGTGCCCTTAACAGCTCCCTGCAGCTCCTG
GAGGACCGTCTGCACCAGCTCAGCCTGAAGGACCTCACTGGGCTGCAGGAGAGGCTGGGGCCCCAGGGCCTCCT
GGGCTGCAGGGACCCCCAGGCCCTGCTGGACCTCAGGATCACCAGGCAAGGACGGGCAAGAGGGCCCCATCGGG
CCACCAGTCTCAAGGGGAACAGGGAGTGGAGGGGGCACCAGGCCCTGTGCCCCAAGTGGCATTTTTCAGCT
GCTCTGAGTTTCCCCGGTCTGAACAGGCACGGTCCCCCTTCGACAGAGTCTGCTCAATGATGGAGGCTATTAT
GATCCAGAGACAGGCGGTGTTACAGCGCCACTGGCTGGACGCTACTTGCTGAGCGCGGTGCTGACTGGGCACCGG
CAGGAGAAAGTGGAGGCCGTGCTGTCCCGCTCCAACAGGCGGTGGCCCGCGTAGACTCCGGTGGCTACGAGCCT
GAGGGCCTGGAGAATAAGCCGGTGGCCGAGAGCCAGCCAGCCGGGACCCCTGGGCGTCTTCAGCCTCATCCTG
CCGCTGCAGGCCGGGGACACGGTCTGCGCTCGACCTGGTTCATGGGGCAGCTGGCGCACTCGGAGGAGCCGCTCACC
ATCTTCAGCGGGGCCCTGCTCTATGGGGACCCAGAGCTTGAACACGCGTAGACTGGGGTCCCCCCCCGACGTGTCT
ACGTGCGGTGAAGAGACAGCGGGGCGGGGCTCTGGGGTCTCGCTGAGACGGGGCACCTAGCCCTGGGCGA
GCGCCGACCCGGGCGCAGCGGCACCGCGCCAGAGCGGCTCTCCCCACGCCCCGGGCGCGCCGGCTCAGGG
AGGCTCGGGGCGCCCATGCAGACTTTTGGCCTGGCGCATCCCCCAAGAACCCTCCAGGGCGGCGCTGCGGAG
GAGCCGATCCTCGCACCCCTCCGCTCCCTCCACTGGCCCTCAGGTGATTCCTGGGCTCCAGGCTCCCCCGCGC
GGGCGCCGCCACCGCCATACTAAACGATCGAGGAATAAAGACACTTGTTTCT

756/6881
FIGURE 706

MAPRTLWSCYLCCLLTAAAGAASYPPRGFSLYTGSSGALSPGGPQAQIAPRPASRHRNWCAVVTTRTVSCVLEDG
VETYVKYQPCAWGQPQCPQSIMYRRFLRPYRVAYKTVTDMEWRCQGYGGDDCAESPAPALGPASSTPRPLARP
ARPNLSGSSAGSPLSGLGGEGPGESEKVQQLLEEQVQSLTKELQGLRGVLQGLSGRLAEDVQRAVETAFNGRQOPA
DAAARPGVHETLNEIQHQLQLLDTRVSTHDQELGHLNNHHGGSSSSSGSRAPAPASAPPGPSEELLRQLEQRLQE
SCSVCLAGLDGFRQQQEDRERLRAMEKLLASVEERQRHLAAGLAVGRRPPQECCSPELGRRLAELERRLDVVAGS
VTVLSGRRGTTELGAAGQGGHPPGYTSLASRLSRLEDNFNSTLGPSEEQEEWPAPGGLSHWLPAAARGRLEQLG
GLLANVSGELGGRDLLEEQVAGAMQACGQLCSGAPGEQDSQVSEILSALERRVLDSEGQLRLVGSGLHTVEAAG
EARQATLEGLQEVVGRQLQDRVDAQDETAEEFTLRNLTAARLGQLEGLLQAHGDEGCGACGGVQEELGRRLRDGVE
RCSCPLLPPRPGAGPGVGGPSRGPLDGFVSFVGSSGSALQALQGESEVILSFSSLNDSLNELQTTVEGQGADL
ADLGATKDRIISEINRLQEQATEHATESEERFRGLEEGQAQAGQCPSLEGRRLGRLEGVCERLDTVAGGLQGLREG
LSRHVAGLWAGLRETNNTSQMQAALLEKLVGGQAGLGRRLGALNSSQLLEDRLHQLSLKDLTG PAGEAGPPGPP
GLQGPAGPAGPPGSPGKDGQEGPIGPPGPQGEQGVGAPAAPVPQVAFSAALSLPRSEPGTVVPFDRVLLNDGGYY
DPETGVFTAPLAGRYLLSAVLTGHRHEKVEAVLSRSNQGVARVDSGGYEPEGLENKPVAESQPSPGTILGVFSLIL
PLQAGDTVCDLVMGQLAHSEEPLTIFSGALLYGDPELEHA

757/6881
FIGURE 707

GGCACGAGGGGACCGCGGGCGCTCGGGTGTTTTTGGGGGCCCCGGGTGGAGGGCCCCGGGTGCCGGGGCCCCAAGGTG
CGGCCTCGCTAGCGGGAGAGGGAGCGGGATCACCGGCCCGGAGAGAGCTCTCAGGGCCAGAGCGGGGCAGGAGGA
TGCTTTCCAGCCCCACCATGGAGCTGCGCTGTGGGGGATTGCTGTTCACTTCTCGCTTTGATTACAGGAATCTA
GCCACGTGGAGAAGGTGGAATCTTTGTCCAGTGATGGGGAAGGGGTAGGAGGTGGGGCGTCAGCCCTGACCACT
GGCATTGCCTCTTCCCTGACTATGAATTCAACGTGTGGACCCGACCAGACTGTGCTGAAACGGAATTTGAGAAT
GGGAACAGGTCATGGTTCTACTTCAGCGTCCGGGGAGGAATGCCAGGAAACTCATCAAGATCAACATTATGAAC
ATGAACAAGCAGAGCAAGCTGTATTCCAGGGCATGGCCCCCTTTGTGCGCACACTGCCACCCGGGCCACGCTGG
GAACGCATTTCGAGACCGGGCCACCTTTGAGATGACAGAGACGCAGTTTGTTATCCTTTGTTTCATCGTTTCGTG
GAGGGCCGTGGGGCCACCACTTTCTCGCTTCTGCTACCCCTTCTCTACAGTGACTGCCAGGAAGTGTAAAC
CAGCTAGACCAGCGCTTTCCGGAGAACCACCTTACCCATAGCAGCCCCCTGGATACCATCTATTACCATCGGGAG
CTCCTTTGCTATTCTCTGGATGGACTTCGTGTAGATCTGCTGACGATCACTTCTGCCATGGGCTTCGAGAAGAT
CGAGAGCCCCGTCTAGAGCAGCTATTTCCTGATACCAGCACCCTCGACCATTCGTTTCGAGGCAAGAGGATA
TTCTTCTTAAGCAGTAGAGTACACCCAGGGGAGACTCCATCTAGCTTTGTCTTCAATGGCTTCTGGAAGTTCATC
CTCCGACCTGATGATCCCCGGGGCCAAACCCCTCCGTCGCTCTTCTGCTTTAAGCTGATTCCCATGTTGAACCCC
GATGGTGTGGTCCGGGGACACTACCGCACAGACTCACGTGGAGTGAATCTGAACCGTCAGTACCTGAAGCCGTGAT
GCCGTCTGCACCCGGCCATCTATGGGGCCAAAGCTGTGCTTCTTACCACCATGTGCACCTCTCGTCTGAAGTCC
CAGAGTTCTCTGAGCACCAGCCAGTTCCTGTCTCCCTCTGATGCTCTGTTTCTGACCTGGAGAAAGCCAAAC
AATCTCCAAATGAAGCTCAGTGTGGGCACTCAGCTGACAGGCATAACGCTGAAGCCCTGGAAACAAACAGAGCCA
GCAGAACAGAAGCTCAACAGTGTGTGATTATGCCACAACAGTCTGCGGGGCTTGAAGAGTCAGCCCCCTGATACC
ATCCCCCCCCAAGAGAGTGGCGTTGCTTACTATGTGGACCTGCATGGACATGCTTCCAAAGGGGCTGCTTCATG
TACGGAAACAGCTTTAGTGATGAGAGCACCAGGTGGAAACATGCTATATCCAAAGCTCATCTCCTTGAATTCA
GCCCACTTCGACTTCCAGGGCTGCAATTTCTCAGAGAAGAATATGTATGCCCGAGACCGTAGAGATGGCCAGTCT
AAAGAGGGAAGCGGCCGTGTTGCAATCTACAAAGCCTCAGGGATAATCCACAGCTACACACTTGAATGCAACTAC
AACACTGGACGCTCAGTAAACAGCATCCCTGCTGCCTGCCATGACAATGGGCGTGCCAGCCCCCTCCCCCGCCG
GCTTTCCCTCCAGATACACTGTGGAATATTGTAGCAGTGGGACGAGCTATGGCCATTGCAGCCCTGGACATG
GCGGAATGTAATCCGTGGCCCCGAATTGTACTGTGAGCAGCAGCAGCCTTACTAATCTACGGGCTGGATGCTG
AAACATGTACGCAACAGCCGAGGCTAAGCAGCACTCTGAATGTGGGTGTCAACAAGAAGAGGGGCTTTCGAAGT
CCACCCAAAAGTCAAAATGGGTTGCCGTCTCCTGCTCCGAAAACACCTTGAGTCGGGCACGAAGTTTTAGCACC
GGCACAAGTCCGGTGTGTAGCAGCAGCAGCCAAACAAATTTCTCCACAGATGAAGAATTCCCCAGCTTTCTTTT
CATGGCAGTCGGCCTGCAGGGCTGCCAGGCTTGGGCTCTAGTACCCAAAAGGTCACCCACCGGGTGCTGGGCCCC
GTCAGAGAGCCCCGAAGCCAGGACAGGAGACGGCAGCAGCAGCAGCCTGAACCATCGTCTGACAGGAGCCTCGCT
CCATCCCCAGCTCTCTAGTTCTGGCCAGCCTCCTCACACAAGCTGGGCTCCTGTCTACTGCTGATTTCATTC
AACATACCAGGGAGCGATTGCTCACTCTTGTCTCTGGAGACAAACCAGAGGCTGTGATGGTAATCGGGAAAGGT
CTGCTAGGGACTGGAGCTCGGATGCCCTGCATCAAGACTCGATTGACAGCCTGTCCGAGGAGAGTTTCCGCCAGG
AGGGGTCCCGGATTCCCCAGGCTAGGCCAGGTTGGGCGGGGCTCACCGCCGACTCGCAGAGGGATGAAGGCT
CTTACAGGCCCCACATCCCCCTACCCCCCGGACCAGGGAGAGCAGTGAGCTGGAGCTGGGATCCTGCTCTGCTACAC
CAGGGCTGCCTCAGGCCAGGCCCCACGGCCCCGCTCTGCCCTGCCTTTTCTCCTATATCCTGTAGTCTATCTG
ACTCCCCATCCTGGAATTGTTACAGCAGGGGTCCCTTGGGCCAACCTGAGGTTTGTGTTTGTCCCTAAATCTCCCC
CACTGACTGTTTTCTCCCGGGTCTGATAATGCCTTTATGTTCAATCCAGGATATAGCCCCAAGATGGGGTAACA
GTGGGAAATATGCTAGTTCCCTCCAGGCGCTGATTCCATGTGACAGCCGTTAAGTCTTGGAAATGCCAGCCAC
GCTGTCCAAGGCATTACAGAGTATCACCTTGAGACAGAACAAAACAGGGACCTGCCACCCCTTCCCTCCCTCCGC
AGCACAAGATTTTGGGACCACAAAAAAGTCTATATTTTATATTGGGGGGAGGGAGTAGAAAAGCAAGCCCCCT
ATACTGGGCCCATTTCAGTGGCAGCTTCTTGTTCATAGGATTAAGGAAGACTCTGAGGAAATAAAAGTTGTTTG
GAAAAATCCAAAAAATAAAAAAAAAA

758/6881

FIGURE 708

MNMNKQSKLYSQGMAPFVRTLPTRPRWERIRDRPTFEMTETQFVLSFVHRFVEGRGATTFFAFICYPFSYSDCQEL
LNQLDQRFENHPHTSSPLDTIYYHRELLCYSLDGLRVDLLTITSCHGLREDREPRLEQLFPDTSTPRPFRFAGK
RIFFLSSRVHPGETPSSFFVNGFLDFILRPDDPRAQTLRRLFVFKLIPMLNPDGVVRGHYRTDSRGVNLNRQYLK
PDAVLHPAIYGAKAVLLYHHVHSRLNSQSSSEHQPSCLPPDAPVSDLEKANNLQNEAQCCHSADRHNAAEWKQT
EPAEQKLSNVWIMPQQSAGLEESAPDTIPPKESGVAYYVDLHGASKRGCFMYGNSFSDESTQVENMLYPKLISL
NSAHFDFQGCNFSEKNMYARDRRDQGSKEGSGRVAIYKASGIIHSYTLNNTGRSVNSIPAACHDNGRASPPP
PPAFPSRYTVELFEQVGRAMAIAALDMAECNPWPRIVLSEHSSLTNLRAWMLKHVRNSRGLSSTLNVGVNKKRGL
RTPPKSHNGLPVSCSENTLSRARSFSTGTSAGGSSSSQONSPQMKNPSFPFHGSRPAGLPGLGSSTQKVTHRVL
GPVREPRSQDRRRQQQPLNHRPAGSLAPSPAPTSSGPASSHKLGSCLLPDSFNIPGSSCSLLSSGDKPEAVMVIG
KGLLGTGARMPCIKTRLQTCPRRVSARRGPGFPRLGPGWAGAHRRRLAEG

759/6881
FIGURE 709

CCCCCCCCACTCTGGACTCCCGCGCTGGGCGCGCTGAGGCGGCCCCCGAGCGAGCGCGCTGCAGCCGCCGCCG
CCCCGAGCACC CGCAGCTCCGGCGCCGCGGCGAGACGGAGACGGACCGAGCCACGGGCCCCCGCGGCCGAGCATC
TCGGAGGAGAACATGCTTGCCAACTCAGCCAGCGTGAGGATCCTCATCAAGGGAGGCAAGGTGGTGAACGATGAC
TGACCCACGAGGCTGACGTCTACATCGAGAATGGCATCATCCAGCAGGTGGGCGCGAGCTCATGATCCCTGGC
GGGGCCAAGGTGATTGATGCCACAGGAAAAGTGGTGATCCCTGGTGGCATCGACACCAGCACCACCTTCCACCAG
ACCTTCATGAATGCCACGTGCGTGGACGACTTCTACCATGGGACCAAGGCAGCACTCGTCGGAGGCACCACCATG
ATCATCGGCCACGTCTGCCCCGACAAGGAGACCTCCCTTGTGGACGCTTATGAGAAGTGGCAGGTCTGGCCGAC
CCCAAGGTCTGCTGTGATTACGCCCTCCACGTGGGGATCACCTGGTGGGCACCCAAGGTGAAAGCAGAAATGGAG
ACACTGGTGAGGGAGAAGGGTGTCACTCGTTCCAGATGTTTCATGACCTACAAGGACCTGTACATGCTTCGAGAC
AGTGAGCTGTACCAAGTGTTCACGCTTGCAAGGACATTGGGGCAATCGCCCGCTCCATGTGAAAATGGGGAG
CTTGTTGGCCGAGGGTGCTAAGGAGGCACTGGATTGGGGATCACAGGCCCAGAAGGAATCGAGATCAGCCGTCCA
GAGGAGCTGGAAGCTGAAGCCACTCATCGTGTATCACCATTGCAAACAGGACTCACTGTCCAATCTACCTGGT
AACGTGTCCAGTATCTCGGCTGGTGACGTTATCGCAGCTGCTAAGATGCAAGGGAAGGTTGTGCTGGCGGAGACC
ACCACTGCACATGCCACGCTGACAGGCTTACACTACTACCACCAGGACTGGTCCCACGCGGCTGCCTATGTACG
GTGCTTCCCTGAGACTGGACACCAACACCTCAACCTACCTCATGAGCCTGCTGGCCAATGACACTCTGAACATC
GTGGCATCAGATACCCGGCTTTCACCACAAAGCAGAAAGCTATGGGCAAGGAAGACTTCACCAAGATCCACAT
GGAGTGAGTGGCGTGACGAGCCGATGAGCGTCATCTGGGAGAGAGGAGTGGTGGAGGAAAGATGGATGAGAAC
CGTTTTGTGGCCGTTACAGTTCACACGAGCTAAGCTTCTGAACCTGTATCCCCGCAAGGGCCGATTATTCCC
GGAGCCGATGCTGATGTGGTGGTGTGGGACCCAGAAGCCACAAAGACCATCTCAGCCAGCACGCAGGTCCAGGGA
GGAGACTTCAACCTGTATGAGAACATGCGCTGCCACGGCGTGCCACTGGTCACCATCAGCCGGGGCGCGTCTGT
TATGAGAACGGCGTCTTCATGTGCGCCGAGGGCACCGGCAAGTCTGTCTCCCTGAGGTCTTCCCAGACACTGTC
TACAAGAAGCTGGTCCAGAGAGAGAAGACTTTAAAGGTTAGAGGAGTGAGCCGACTCCCTACCTGGGGGATGTC
GCTGTTGTGCTGCACCTGGGAAAAAAGAGATGGGAACCCCACTCGCAGACACTCCTACCCGGCCCGTCAACCGG
CATGGGGGATGAGGGACCTTACGAATCCAGCTTCAGCCTCTCTGGCTCTCAGATCGATGACCATGTTCCAAAG
CGAGCTTCAGCTCGGATCCTCGCTCCTCCCGGAGGCAGGTGAGTGGCATTGTTGGTAAAGGCATTGCCAAGCCCC
CGAGTGAGGACGCACCGCCGCCACAGCCGCAACTCTCCAGCCGAAGCTGCAGGGGCAGGAGAGGCTGGGCTGG
GTGGCACACCACCCGAGGGGGGGCCCCGGGACCCACGGAGCCCTCCCTATGTCTGCAAAGTGATTCACTGTGCTTC
GAGCCAACTCTAACAGGCACCTTGAGATGTGTTCCCTCCTGCTGTAGTCCCTTCTGCTTGGCCTCGGCGGGCTTT
TCTGGGGCCCGAGGAAGCCACACTATGCACAGAGCCCAATGCATAGAGCCCTGGCCAGCCCTTCTCTCACTCCT
GCCTCCGCTGGCTTTGGGAAAGCCAGACTTTAGTGCCCTGCCCTTGGCTGACTGGCCAGTTGCCAGAGCACT
TTAGCAGATGTGGTTTCAAAGTAAAGGCCTCTCCCCACCCCTTAGGCCCCGTGGTGACATTTCCCAAGTCAGA
CAGATGTCAGCTTCCCAGCCATGCCAGGACGTCTATCTCCCCAACCCACCTCTGGCCCTGTGTAGGGGCAGG
GATGGGGGTGGCTGGGACTCCTGGTGGCCCTCGCCAGCTTCTCCTGCGCCCCGCCCCACCCCTCGGGGGGTGAC
AGGCCCAGAAGGGTAGCTGGGCGGGGCTCGAGGCTGGTGCCAGGCGCGTGTAATGGTTTTGTTTTGCACGTTG
GTTTTGCGCAGTAGTTTGGTTTGACTTGTGTTGTGTCATCCTGTGAAAAATAACGGTGCTTGTGTCACTAGCATAGAA
TAGCGACAGGAATAGATGTGGTCTTAGGAGACGTGCACCTTGACACCAACCAGACAGCACAGGGCAGGGGTGGT
GGAGGGGGCTGGGCTCACAGGCCTCTCTTTTCCCGCCTGCAGTCTTCTGGGCTGCGGGAGGCCCTGGCCCTTTC
CCCTTCCCTCCCTCCTTGTCTAGTTTCCACATTCCAAAAGGGGGCTGGGATGCTAGCCCCAGAGATGCCAG
CCCTTCAGGAAGCAGGTGTCTTTCCCTCTCTGCCCTGATCACTCCCAGCACTCCCTTGCCTTCCCTGTCT
TCACCTGCCACCACACACACACACACACACACACACACACACATGGCTTCTATAACTTCTTCTGCTG
GACAGAGACTCAGCGCTCCTCCTGTGTGACTGGCAAGAGGCCTCATGCCTGCTGAGAGAGGGTTCGACGGGCGG

760/6881

FIGURE 710

MLANSASVRILIKGGKVVNDCTHEADVYIENGIIQQVGRELMIPGGAKVIDATGKLVIPGGIDTSTHFHQTFMN
ATCVDDFYHGTKAALVGGTTMIIGHVLPDKETSLVDAYEKCRGLADPKVCCDYALHVGITWWAPKVKAEMETLVR
EKGVNSFQMFMTYKDLYMLRDSELYQVLHACKDIGAIARVHAENGELVAEGAKEALDLGITGPEGIEISRPEELE
AEATHRVITIANRTHCPIYLVNVSSISAGDVIAAAKMQGKVLAETTTAHATLTGLHYHQDWSHAAAYVTVPPL
RLDTNTSTYLMSELLANDTLNIVASDHRPFTTKQKAMKEDFTKIPHGVSGVQDRMSVIWERGTVGGKMDENRFVA
VTSSNAKLLNLYPRKGRIIPGADADVVDPEATKTISASTQVGGDFNLYENMRCHGVPLVTISRGRVYENG
VFMCAEGTGKFCPLRSFPDVTYKKLVQREKTLKVRGVDRTPYLGDVAVVVHHPGKKEMGTPLADTPTRPVTRHGM
RDLHESSFSLSGSQIDDHVPKRASARILAPPGGRSSGIW

761/6881
FIGURE 711

CTTGCTCCGAGAGGGAGTCCTCGCGGACGTCAGCCAAGATTCCAGAAATGACTATCTTGA¹CTTACCCTTTAAAAA
TCTTCCCAGTGCATCAAAATGGGCCCTCAGATTTTCCATAAGACCTCTGAGCTGTTCCCTCCCAGCTACGAGCTGC
CCCAGCTGTCCAGACCAAAACGAAGAAGACGTTAGCCAAACCAATATAAGGAATGTTGTGGTGGTGGATGGTGT
TCGCACTCCATTTTTGTGTCTGGCACTTCATATAAAGACCTGATGCCACATGATTTGGCTAGAGCAGCGCTTAC
GGGTTTGTGTGCATCGGACCAAGTGTCCCTAAGGAAGTAGTTGATTATATCATCTTTGGTACAGTTATTCAGGAAGT
GAAAACAAGCAATGTGGCTAGAGAGGCTGCCCTTGGAGCTGGCTTCTCTGACAAGACTCCTGCTCACACTGTCAC
CATGGCTTGTATCTCTGCCAACCAAGCCATGACCACAGGTGTTGGCTTGATTGCTTCTGGCCAGTGTGATGTGAT
CGTGGCAGGTGGTGTGAGTTGATGTCCGATGTCCCTATTTCGTCACTCAAGGAAAATGAGAAAATGATGCTTGA
TCTCAATAAGGCCAAATCTATGGGCCAGCGACTGTCTTTAATCTCTAAATTCCGATTTAATTTCCCTAGCACCTGA
GCTCCCTGCGGTTTCTGAGTTCTCCACCAAGTGAAGCCATGGGCCACTCTGCAGACCGACTGGCCGCTGCCTTTGC
TGTTTCTCGGCTGGAAACAGGATGAATATGCACTGCGCTCTCACAGTCTAGCCAAGAAGGCACAGGATGAAGGACT
CCTTCTGATGTGGTACCCTTCAAAGTACCAGGAAAAGATACAGTTACCAAAGATAATGGCATCCGCTCCTTCCTC
ACTGGAGCAGATGGCCAAACTAAAACCTGCATTATCAAGCCCTACGGCACAGTGACAGCTGCAAATTCCTTCTT
CTTGA²CTGATGGTGCATCTGCAATGTTAATCATGGCGGAGGAAAAGGCTCTGGCCATGGGTTATAAGCCGAAGGC
ATATTTGAGGGATTTTATGTATGTGCTCAGGATCCAAAGATCAACTATTACTTGGACCAACATATGCTACTCC
AAAAGTTCTAGAAAAGGCAGGATTGACCATGAATGATATTGATGCTTTTGAATTTTATGAAGCTTCTCGGGTCA
GATTTTGGCAAATTTTAAAGCCATGGATTCTGATTGGTTTGCAGAAAATACATGGGTAGAAAAACCAAGGTTGG
ATTGCCTCCTTTGGAGAAGTTTAAATAACTGGGGTGGATCTCTGTCCCTGGGACACCCATTTGGAGCCACTGGCTG
CAGGTTGGTCATGGCTGCTGCCAACAGATTACGGAAAGAAGGAGGCCAGTATGGCTTAGTGGCTGCGTGTGCAGC
TGGAGGGCAGGGCCATGCTATGATAGTGGAGCTTATCCAAAAT³TAATAGATCCAGAAGAAGTGACCTGAAGTTTC
TGTGCAACACTCACACTAGGCAATGCCATTCAATGCATTACTAAATGACATTGTAGTTCCCTAGCTCCTCTTAG
GAAAACAGTTCTTGTGGCCTTCTATTAAATAGTTTGCACCTTAAGCCTTGCCAGTGTCTGAGCTTTTCAATAATC
AGTTTACTGCTCTTTCAGGGATTTCTAAGCCACCAGAATCTCATGAGATGTGTGGGTGGTTGTTTTTGGTCTC
TGTTGTCACTAAAGACTAAATGAGGGTTTGCAGTTGGGAAAGAGGTCAACTGAGATTGGAAATCATCTTTGTAA
TATTTGCAAATTATACTTGTCTTATCTGTGTCCTAAAGATGTGTTCTCTATAAAATACAAACCAACGTGCCTAA
TTAATTATGGAAAAATAATTCAGAATCTAAACACCACTGAAAACCTTATAAAAAATGTTTAGATACATAAATATGG
TGGTCAGCGTTAATAAAGTGGAGAAATATTGGAAAAA

762/6881

FIGURE 712

MTILTYPFKNLPTASKWALRFSIRPLSCSSQLRAAPAVQTKKTLAKPNIRNVVVVDGVRTPFLLSGTSYKDLM
PHDLARAALTGLLHRTSVPKEVVYIIFGTVIQEVKTSNVAREEALGAGFSDKTPAHTVTMACISANQAMTTGVG
LIASGQCDVIVAGGVELMSDVP IRHSRKMRKMLDLNKAQSMGQRLSLISKFRFNFLAPELPAVSEFSTSETMGH
SADRLAAAFVSRLEQDEYALRSHSLAKKAQDEGLSDVVPFKVPGKDTVTKDNGIRPSSLEQMAKLPKPAFIKPY
GTVTAANSSFLTGDASAMLIMAEKALAMGYKPKAYLRDFMYVSQDPKDQLLLGPTYATPKVLEKAGLTMNDIDA
FEFHEAFSGQILANFKAMDSWFAENYMGRKTKVGLPPELFNNWGGSLSLGHPFGATGCRLVMAAANRLRKEGG
QYGLVAACAAGGQGHAMIVEAYPK

763/6881
FIGURE 713

GGCACGAGGCCACTGCTGTCTCTTCAGCTCAAGATGGTGGCCTGCCGGGCGATTGGCATCCTCAGCCGCTTTTC
TGCCTTCAGGATCCTCCGCTCCCGAGGTTATATATGCCGCAATTTTACAGGGTCTTCTGCTTTGCTGACCAGAAC
CCATATTAACATATGGAGTCAAAGGGGATGTGGCAGTTGTTTGAATTAACCTCTCCCAATTCAAAGGTAAATACACT
GAGTAAAGAGCTACATTCAGAGTTCTCAGAAGTTATGAATGAAATCTGGGCTAGTGATCAAATCAGAAGTGCCGT
CCTTATCTCATCAAAGCCAGGCTGCTTTATTGCAGGTGCTGATATCAACATGTTAGCCGCTTGCAAGACCCTTCA
AGAAGTAACACAGCTATCACAAGAAGCACAGAGAATAGTTGAGAACTTGAAAAGTCCACAAAGCCTATTGTGGC
TGCCATCAATGGATCCTGCCTGGGAGGAGGACTTGAGGTTGCCATTTTCATGCCAATACAGAATAGCAACAAAAGA
CAGAAAAACAGTATTAGGTACCCCTGAAGTTTTGCTGGGGGCCCTTACCAGGAGCAGGAGGCACACAAAGGCTGCC
CAAAATGGTGGGTGTGCCTGCTGCTTTGGACATGATGCTGACTGGTAGAAGCATTTCGTGCAGACAGGGCAAAGAA
AATGGGACTGGTTGACCAACTGGTGGAAACCCCTGGGACCAGGACTAAAACCTCCAGAGGAACGGACAATAGAATA
CCTAGAAGAAGTTGCAATTACTTTTGCCAAAGGACTAGCTGATAAGAAGATCTCTCCAAAGAGAGACAAGGGATT
GGTGGAAAAATTGACAGCGTATGCCATGACTATTCCATTTGTCAGGCAACAGGTTTACAAAAAGTGGAAGAAAA
AGTGCGAAAGCAGACTAAAGGCCCTTTATCCTGCACCTCTGAAAATAATTGATGTGGTAAAGACTGGAATTGAGCA
AGGGAGTGATGCCGTTATCTCTGTGAATCTCAGAAATTTGGAGAGCTTGAATGACCAAGAATCAAAGGCCCTT
GATGGGACTCTACCATGGTCAGGTCTGTGCAAGAAGAATAAATTTGGAGCTCCACAGAAGGATGTTAAGCATCT
GGCTATTCTTGGTGACGGGCTGATGGGAGCAGGCATCGCCCAAGTCTCCGTGGATAAGGGGCTAAAGACTATACT
TAAAGATGCCACCTCACTGCGCTAGACCGAGGACAGCAACAAGTGTTCAAAGGATTGAATGACAAAGTGAAGAA
GAAAGCTCTAACATCATTTGAAAGGGATTCCATCTTCAGCAACTTGACTGGGCAGCTTGATTACCAAGGTTTTGA
AAAGGCCGACATGGTGATTGAAGCTGTGTTTGAGGACCTTAGTCTTAAGCACAGAGTGCTAAAGGAAGTAGAAGC
GGTGATTCCAGATCACTGTATCTTTGCCAGTAACACATCTGCTCTCCCAATCAGTGAATCGCTGCTGTCTAGCAA
AAGACCTGAGAAGGTGATTGGCATGCACTACTTCTCTCCCGTGGACAAGATGCAGCTGCTGGAGATTATCACGAC
CGAGAAAACCTTCCAAAGACACCAAGTGCTTCAGCTGTAGCAGTTGGTCTCAAGCAGGGGAAGGTCATCATTGTGGT
TAAGGATGGACCTGGCTTCTATACTACCAGGTGTCTTGCGCCCATGATGTCTGAAGTCATCCGAATCCTCCAGGA
AGGAGTTGACCCGAAGAAGCTGGATTCCCTGACCACAAGCTTTGGCTTTCCCTGTGGGTGCCGCCACACTGGTGGA
TGAAGTTGGTGTGGATGTAGCGAAACATGTGGCGGAAGATCTGGGCAAAGTCTTTGGGGAGCGGTTTGGAGGTGG
AAACCCAGAAGCTGCTGACACAGATGGTGTCCAAGGGCTTCCTAGGTCGTAAATCTGGGAAGGGCTTTTACATCTA
TCAGGAGGGTGTGAAGAGGAAGGATTTGAATCTGACATGGATAGTATTTTAGCGAGTCTGAAGCTGCCTCCTAA
GTCTGAAGTCTCATCAGACGAAGACATCCAGTTCCGCCTGGTGACAAGATTGTGAATGAGGCAGTCATGTGCCT
GCAAGAGGGGATCTTGCCACACCTGCAGAGGGAGACATCGGAGCCGTCTTTGGGCTTGGCTTCCCGCCTTGCTCT
GGGAGGGCCTTTCCGCTTTGTGGATCTGTATGGCGCCAGAAGATAGTGGACCGGCTCAAGAAATATGAAGCTGC
CTATGGAAAACAGTTACCCCATGCCAGCTGCTAGCTGACCTAGCTAACAGCCCTAACAAAGAAGTTCTACCAGTG
AGCAGGCCTCATGCCTCGCTCAGTCAGTGCACCTAACCCAGCTGCCGGCAGTGTGTTCTTCCAACAGAGTGGTG
TCTAGATTTATCAGAGTAACGAGAAGACAAACTCCGGCACTGGGTTTGCTCCCTGATTAAAGTGCCTTCAGCCAA
GACCATCTCTCCCTCCTGGTGAAGTGTGACTTCGAATTAGTTTGACTTCTTATTGGAAGGTAGAGCCCACTGCT
CATGTGATAAGCCCCGAGGCCTAGAGTGGCAGCCAAGAGCCATCTGAAGCCACCTCTCTGCCTGTTTCTCCCAAG
AGGCCAGGGTGGCCAGGGGTGGTGAGGGCAGTTCTGCACCCAGCCAAACACATAACAATAAAAAACCAAACCTCTGT
GTCAGCATCTTTGCCCTTCTGGTTTTAAACGCCTCCTTCAAAAAGCAATCTGGAAGAAAGCCCTGTGCTTTGGGGG
AGTAAGAATGTGTGTGCAATTTCTAGGCAGCACCTTAGGGAGGGACTGGGATGAGAGAAAGTGGGACCTGGTGG
GCTCAACCACACACCTGTCTGTGCAAGTCTTTGCCAGGCTTCTCACCACGGTGTACCGGGATATTAAACCT
CTTCCCCAGCAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA

764/6881
FIGURE 714

MVACRAIGILSRFSAFRILRSRGYICRNFTGSSALLTRTHINYGVKGDVAVVRINSPNSKVNTLSKELHSEFSEV
MNEIWASDQIRSAVLISSEKPGCFIAGADINMLAACKTLQEVTLQSQAQRIVEKLEKSTKPIVAAINGSCLGGL
EVAISCQYRIATKDRKTVLGTPEVLLGALPGAGGTQRLPKMVGVPAAALDMLTGRSIRADRAKKMGLVDQLVEPL
GPGLKPPEERTIEYLEEVAITFAKGLADKKISPKRDKGLVEKLTAYAMTIPFVRQQVYKKVEEKVRKQTKGLYP
PLKIIDVVKTGIEQGS DAGYLCESQKFGE LVMTKESKALMGLYHGQVLCCKNKFGAPQKDVKHLAILGAGLMGAG
IAQVSVDKGLKTI LKDATLTALDRGQQQVFKGLNDKVKKKALTSFERDSIFS NLTGQLDYQGF EKADMVIEAVFE
DLSLKHRVLKEVEA VIPDHCIFASNTSALP ISEIAAVSKRPEK VIGMHYFSPVDKMQLLEIITTEKTSKDT S
VAVGLKQGVIIIVKDGPGFYTT RCLAPMMSEVIRILQEGVDPKKLDSLTT SFGFPVGAATLVDEVGVDVAKHVA
EDLGKVFGERF GGNPELLTQMVSKGFLGRKSGKGFYIYQEGVKRKDLNSDMSILASLKLPPKSEVSSDEDIQF
RLVTRFVNEAVMCLQEGILATPAEGDIGAVFGLGFPCLGGPFRFVDLYGAQKIVDRLKKYEAA YGKQFTPCQLL
ADHANSPNKKFYQ

765/6881

FIGURE 715

TTCTGTTTTTCTCTCTCATTCTCCAGTGGCGGCGGGGGAAGGCGGAGGCAGAGGCAGCAGCAGCCGCGCTGGC
TGCAATGAATGATCCCCAGCTTGGGGGGAGGACTCCAGGTGAGCCTCTGCCCTCGGGAGGCCCGGGACCCCGG
CCGCCCACGACCGGCAGCCACGCTATGGATCCCTAGAGGAAGGAGGAGAAGACAGCTCGCCGCCACCCCATC
CCATTTTCCTCTTCCTTTATCTCATTGTTGCCGAAGCTGTTTACGGCAGCGCTCCCTCTGCTCCAGGACTCCCTG
GGGGGAATGCCAAGACCATCATGGTAGCCACACTGGGGCCAGCTTCTCACAGCTACGATGAGAGCCTCTCCACC
TTGCGCTTTGCCAACCGAGCCAAGAACATCAAGAACAAGCCCCAGGTGAACGAGGACCCCAAGGACACACTGCTG
CGGGAATTCCAAGAGGAGATTGCCCCGCTGAAGGCCCAGCTGGAGAAGAGGGGGATGCTGGGGAAGCGGCCCCGG
AGGAAGAGCAGCCGCAGGAAGAAGGCCGTGTCCGCCCCGCTGGGTACCCTGAGGGCCCAGTGATTGAGGCCTGG
GTGGCAGAAGAGGAGGATGACAACAACAACCAACCGCCCGCCCCAGCCATCCTGGAGTCAGCCTTGGAGAAG
AACATGGAGAATTACCTGCAGGAACAGAAGGAGCGGCTGGAGGAGGAGAAGGCAGCCATCCAGGATGACCGCA

766/6881
FIGURE 716A

GCTGGAGGTGGCCTCCCCTCCGCCCCAGACAAGAAGAGGCCCTCAGCCCTCCCCGGTCTCAGAGAGCCCTGAGA
GGAGGCCAGTCCAGAGCTCTTCTCCGTTCCAGTCCACTTCTCTAGGGCCAGTAGCAGACACCAGCCAGTATG
CCGAGGAACCAGGGCTTCTCCGAGCCCGAATACTCGGCCGAGTACTCAGCCGAGTACTCCGTCAGCCTGCCCTCC
GACCCTGACCGCGGGGTGGGCCGACCCATGAAATCTCGGTCCGGAACTCGGGCTCCTGCCTGTGCCTGCCTCGC
TTCATGCGGTGACTTTTCGTGCCGGAGTCTTGGAGAACCTCTACCAGACCTACTTCAAAGGCAGCGCCACGAG
ACCCTGCTGGTGTGGTGGTCTTTCAGCCCTCTTGGACTGCTACGTGGTGGTTCATGTGTGCTGTGGTCTTCTCC
AGCGACAAGCTGGCTTCCCTCGCCGTGGCTGGAATTGGACTGGTGTGGACATCATCCTCTTCGTGCTCTGCAAA
AAGGGGCTGCTCCCGGACCGGGTCACCCGCAGAGTGTGCTGCTGCTGTGGCTGCTCATAACCGCCAGATC
TTCTCCTACCTGGGCTGAACCTCGCGCGTGCCACGCGGCTAGTGACACGGTGGGCTGGCAGGTCTTCTTTGTC
TTCTCCTTCTTTCATCAGCTGCCCTCAGCCTCAGCCCCATCGTGATCATCTCCGTGGTCTCCTGTGTGGTGCAC
ACGTTGGTCTTGGGGGTACCGTGCGCCAGCAGCAGCAGGAGGAGCTCAAGGGGATGCAGCTGCTGCGGGAGATC
CTGGCCAACGTCTTCTCTACCTGTGCGCCATCGCTGTGGGCATCATGTCTACTACATGGCTGACCGCAAGCAC
CGCAAGGCCTTCTGGAGGCGCCGAGTCTGCTGGAGGTGAAGATGAACCTGGAAGAGCAGAGCCAGCAGCAGGAG
AACCTCATGCTTTCCATCCTGCCCAAGCACGTGGCTGACGAGATGCTGAAAGACATGAAGAAAGACGAGAGCCAG
AAGGACCAGCAGCAGTTC AACACCATGTACATGTACCGTACAGAGAACGTGAGCATCCTCTTTGCCGACATCGTG
GGCTTTACCCAGCTGTCTTCTGCCTGCAGTGCCAGGAGCTTGTGAAGCTGCTCAACGAGCTCTTTGCCCGCTTT
GACAAGCTGGCAGCTAAATACCACAGCTGCGGATTAAGATCCTGGGCGACTGCTACTACTGCATCTGCGGCTTG
CCCGACTACCGGGAGGACCACGCCGTCTGCTCCATCCTCATGGGGCTGGCCATGGTGGAGGCCATCTCGTATGTG
CGGGAGAAGACCAAGACTGGGGTGGACATGCGTGTGGGGGTGCACACGGGCACCGTGTGGGGGGCGTCTGGGC
CAGAAGCGCTGGCAGTACGACGTGTGGTTCGACTGATGTCACTGTAGCCAACAAGATGGAGGCCGCGGCATCCCT
GGGCGCGTGCACATCTCCAGAGCACCATGGACTGCCTGAAAGGGGAGTTTGATGTGGAGCCAGGCGATGGGGC
AGCCGCTGTGATTACCTAGAAGAGAAGGGTATTGAAACCTACCTCATCATTGCCTCCAAGCCAGAGGTGAAGAAA
ACAGCCACCCAGAATGGCTCAATGGCTCGGCCCTGCCCAATGGAGCACCAGCTTCTCAAAGTCCAGCTCCCT
GCCCTCATTGAGACCAAGGAGCCCAACGGGAGTGCCACAGCAGTGGGTCCACGTCCGAGAAGCCCGAGGAGCAG
GATGCCCAGGCCGACAACCCCTCATTCCCCAACCCACGCCGGAGGCTGCGCTGCAGGACCTGGCTGACCGAGTG
GTGGATGCCTCTGAAGATGAGCACGAGCTCAACCAGCTGCTCAACGAGGCCCTGCTTGAGCGAGAGTCCGCCCAA
GTGTAAGAAGAGAAACACCTTCTTGTCCATGCGGTTTCATGGACCCGAGATGGAACCCGCTACTCGGTGG
AGAAGGAGAAGCAGAGTGGGGCTGCCTTCAGCTGCTCCTGCGTCTGCTCTGCACGGCCCTGGTTCGAGATAC
TCATCGACCCCTGGCTAATGACAACTATGTGACCTTCATGGTGGGGGAGATTCTGCTCCTCATCCTGACCATCT
GCTCCCTGGCTGCCATCTTTCCCCGGGCCTTCTCAAGAAGCTGTGGCCTTCTCAACTTGGATTGACCGGACCC
GCTGGGCCAGGAACACCTGGGCCATGCTCGCCATCTTCATCCTGGTGATGGCAAATGTCGTGGACATGCTCAGCT
GTCTCCAGTACTACAGGGACCCAGCAATGCAACGGCAGGGATGGAACGGAGGGCAGCTGCCTGGAGAACCCCA
AGTATTACAACTATGTGGCCGTGCTGTCCCTCATCGCCACCATCATGCTGGTGCAGGTGAGCCACATGGTGAAGC
TCACGCTCATGCTGCTCGTCGAGGCGCCGTGGCCACCATCAACCTCTATGCCTGGCGTCCCGTCTTTGATGAAT
ACGACCACAAGCGTTTTTCGGGAGCACGACTTACCTATGGTGGCCTTAGAGCAGATGCAAGGATTCAACCCCTGGGC
TCAATGGCACTGACAGGCTGCCCCCTGGTGCCTTCCAAGTACTCTATGACGGTGTGGTGTTCCTCATGATGCTCA
GCTTCTACTACTTCTCCGCCACGTAGAAAACTGGCACGGACACTTTTCTTGTGGAAGATTGAGGTCCACGACC
AGAAGGAACGTGTCTATGAGATGCGACGCTGGAACGAGGCCTTGGTCACCAACATGTTGCCTGAGCAGTGGCAC
GCCATTTCTTGGGGTCCAAGAAGAGAGATGAGGAGCTGTATAGCCAGAGTATGATGAGATTGGAGTCATGTTTG
CCTCCCTGCCCACTTTGTGACTTCTACACAGAGGAGCATCAACAATGGTGGTATTGAGTGTCTGCGTTCCT
TCAATGAAATCATCTCAGATTTTGACTCTCTCTGGACAATCCCAAGTTCCGGGTGATCACCAAGATCAAAACCA
TTGGCAGCACGTATATGGCGGCTTCAGGAGTCAACCCCGATGTCAACACCAATGGCTTTGCCAGCTCCAACAAGG
AAGACAAGTCCGAGAGAGAGCGCTGGCAGCACCTGGCTGACCTGGCCGACTTTCGCGCTGGCCATGAAGGATACGC
TCACCAACATCAACAACAGTCTTCAATAACTTCATGCTGCGCATAGGCATGAACAAAGCGGGGTTCTGGCTG
GGGTATCGGAGCCCGAAACCACTACGACATCTGGGGCAATACAGTCAATGTAGCCAGCAGGATGGAGTCCA
CGGGGGTCATGGGCAACATTGAGGTATGTCCAGTGGCACAGCTGGCACGTGCTCAGACTCGGCATGAGAACAAAA
CCGGAACAGTGCTTTCCACGTGCCCCCTGCCCTCCACCCCTGAACAGTAAGTCCAGAGGCAGAGACGTGG
GGCAAGTGGAAAAGAATCTCTGCAATAGCTAGCCTCAGCACAGGAAGAAACACTTCCACCCATCCACAAAATCTG

767/6881

FIGURE 716B

GGCTGAAGTTCTAGATGCATATGGCCTACCTTTGGGTATGAACCTTCTAATACCTTGTGATCTGTCTCAGTGGCG
TCTGGCAGAAAAGCTCCACGAAAGGGGCAAAGGGACAGAAGAGCAGGCAGTGGCAGTTTCCAGGTAACAGTGCTG
CCACCCAGGTGACCATGGCCATCCTAGGAGAGAAGTCTCTGGATGAGAGGAGGCCCCCAGCCAGTCACTCCAGGG
TTCTCCTTCCCACCCGCTGCTCGCCACGGGTGGAAGGGCTTGGCCC

768/6881
FIGURE 717

GCCGCGGCACCAGGGCGCGCAGCCGGGCGGGCCCGACCCACCGGCCATACGGTGGAGCCATCGAAGCCCCACC
CACAGGCTGACAGAGGCACCGTTCCACCAGAGGGCTCAACACCGGGATCTATGTTTAAGTTTAACTCTCGCTCC
AAAGACCACGATAATTCCTTCCCCAAAGCCAGCAGCCCCCAGCCCCGCGCAGCCCCAGCCTGCCTCCCGGCGC
CCAGATGCCCCGCGCATGCCCTCCAGCGGCCCCGGGGACACCAGCAGCTCTGCTGCGGAGCGGGAGGAGACCAGAA
GGACGGAGAGGAGCAGGAGGAGCCGCGTGGCAAGGAGGAGCGCCAAGAGCCCAGCACCACGGCAGCGAAGGTGGG
GCGGCCTGGGAGGAAGCGCAAGCACCCCCCGGTGGAAAGCGGTGACACGCCAAAGGACCCTGCGGTGATCTCCAA
GTCCCCATCCATGGCCAGGACTCAGGCGCTCAGAGCTATTACCCAATGGGGACTTGGAGAAGCGGAGTGAGCC
CCAGCCAGAGGAGGGGAGCCCTGCTGGGGGGCAGAAGGGCGGGGCCCCAGCAGAGGGAGAGGGTGACGCTGAGAC
CCTGCTGAAGCCTCAAGAGCAGTGGAAAATGGCTGCTGCACCCCAAGGAGGGCCGAGGAGCCCTGCAGAAGC
GGGCAAGAAGACAGAAGGAGACCAACATCGAATCCATGAAAATGGAGGGCTCCCGGGGCCGGCTGCGGGGTGGCTT
GGGCTGGGAGTCCAGCCTCCGTCAGCGGCCCATGCCGAGGCTCACCTTCCAGGCGGGGAGCCCTACTACATCAG
CAAGCGCAAGCGGGACGAGTGGCTGGCACGCTGGAAAAGGGAGGCTGAGAAGAAAGCCAAGGTCAATTGCAGGAAT
GAATGCTGTGGAAGAAAACAGGGGCCCGGGGAGTCTCAGAAGGTGGAGGAGGCCAGCCCTCCTGCTGTGACGA
GCCCCACTGACCCCGCATCCCCACTGTGGCTACACGCTGAGCCCGTGGGGTCCGATGCTGGGGACAAGAATGC
CACCAGAGCAGGCGATGACGAGCCAGAGTACGAGGACGGCCGGGGCTTTGGCATTGGGGAGCTGGTGTGGGGAA
ACTGCGGGGCTTCTCCTGGTGGCCAGGCCGCTTGTGTCTTGGTGGATGACGGGCCGAGCCGAGCCAGCTGAAGG
CACCCGCTGGGTGATGTGGTTCCGAGACGGCAAATTCTCAGTGGTGTGTGTTGAGAAGCTGATGCCGCTGAGCTC
GTTTTGCACTGCGTTCCACCAGGCCACGTACAACAAGCAGCCCATGTACCGCAAAGCCATCTACGAGGTCTCTGCA
GGTGGCCAGCAGCCGCGCGGGGAAGCTGTTCCCGGTGTGCCACGACAGCGATGAGAGTGACACTGCCAAGGCCGT
GGAGGTGCAGAACAAGCCCATGATTGAATGGGCCCTGGGGGGCTTCCAGCCTTCTGGCCCTAAGGGCCTGGAGCC
ACCAGAAGAAGAGAAGAATCCCTACAAAGAAGTGACACGGACATGTGGGTGGAACCTGAGGCAGCTGCCTACGC
ACCACCTCCACCAGCCAAAAGCCCCGGAAGAGCACAGCGGAGAAGCCCAAGGTCAAGGAGATTATTGATGAGCG
CACAAGAGAGCGGCTGGTGTACGAGGTGCGGCAGAAGTGCCGGAACATTGAGGACATCTGCATCTCCTGTGGGAG
CCTCAATGTTACCCTGGAACACCCCTCTTCGTTGGAGGAATGTGCCAAAAGTCAAGAACTGCTTTCTGGAGTG
TGCGTACCAGTACGACGACGACGGCTACCACTCTACTGCACCATCTGCTGTGGGGGCGGTGAGGTGCTCATGTG
CGGAAACAACAACTGCTGCAGGTGCTTTTTCGTTGGAGTGTGTGGACCTCTTGGTGGGGCCGGGGGCTGCCAGGC
AGCCATTAAGGAAGACCCCTGGAAGTGTACATGTGCGGGCACAAGGGTACCTACGGGCTGCTGCGGCGGCGAGA
GGACTGGCCCTCCCGGCTCCAGATGTTCTTCGCTAATAACCACGACCAGGAATTTGACCCTCCAAAGGTTTACCC
ACCTGTCCCAGCTGAGAAGAGGAAGCCCATCCGGGTGCTGTCTCTCTTTGATGGAATCGCTACAGGGCTCCTGGT
GCTGAAGGACTTGGGCATTGAGGTGACCGCTACATTGCCTCGGAGGTGTGTGAGGACTCCATCACGGTGGGCAT
GGTGGCGCACAGGGGAAGATCATGTACGTGCGGGACGTCCGCGAGCGTCACACAGAAGCATATCCAGGAGTGGGG
CCCATTCGATCTGGTGATTGGGGGAGTCCCTGCAATGACCTCTCCATCGTCAACCCTGCTCGCAAGGGCCTCTA
CGAGGGCACTGGCCGGCTCTTCTTTGAGTTCTACCGCTCCTGCTATGATGCGCGGCCCAAGGAGGGAGATGATCG
CCCCTTCTTCTGGCTCTTTGAGAATGTGGTGGCCATGGGCGTTAGTGACAAGAGGGACATCTCGCGATTTCTCGA
GTCCAACCCTGTGATGATTGATGCCAAAGAAGTGTGAGCTGCAGGAGTGTCTGGAGCATGGCAGGAT
CGGTATGAACAGGCCGTTGGCATCCACTGTGAATGATAAGCTGGAGCTGCAGGAGTGTCTGGAGCATGGCAGGAT
AGCCAAGTTCAGCAAAGTGAGGACCATTAACGAGGTCAAACCTCCATAAAGCAGGGCAAAGACCAGCATTTTCC
TGCTTTCATGAATGAGAAAGAGGACATCTTATGGTGCATGAAATGGAAAGGGTATTTGGTTTCCAGTCCACTA
TACTGACGTCTCCAACATGAGCCGCTTGGCGAGGCAGAGACTGCTGGGCGGTCATGGAGCGTGCCAGTCATCCG
CCACCTCTTCGCTCCGCTGAAGGAGTATTTTCGCTGTGTGTAAGGACATGGGGGCAAAGTGGGTAGCGACACA
AAGTT

769/6881
FIGURE 718

MPAMPSSGPGDTSSSAAEREEDRKDGEEQEPRGKEERQEPSTTARKVGRPGRKRKHPPVESGDTPKDPAVISKS
PSMAQDSGASELLPNGDLEKRSEPPQEEGSPAGGQKGGAPAEEGEAAETLPEASRAVENGCCTPKEGRGAPAEAG
KEQKETNIESMKMEGSRGRLRGGLGWESSLRQRPMPRLTFQAGDPYYISKRRDEWLARWKREAEKKAKVIAGMN
AVEENQGPGESQKVVEASPPAVQQPTDPASPTVATTPEPVGSDAGDKNATKAGDDEPEYEDGRGFGIGELVWGKL
RGFSWWPGRIVSWWMTGRSRAAEGTRWVMWFGDGKFSVVCVEKLMPLSSFCSAFHQATYNKQPMYRKAIYEVLOV
ASSRAGKLFVPVCHDSDESDTAKAVEVQNKPMIEWALGGFQPSGPKGLEPPEEEKNPYKEYYTDMMWEPEAAAYAP
PPPAKKPRKSTAEKPKVKEIIDERTRERLVYEVQRKCRNIEDICISCGSLNVTLEHPLFVGGMQCNCKNCFLECA
YQYDDDGYSYCTICCGGREVLNCGNNCCRCFCVECDLLVGPAAQAAIKEDPWNCYMCCHKGTGTYGLRRRED
WPSRLQMFFANNHDQEFDPKVPVPAEKRKPIRVLSLFDGIATGLLVKDLGIQVDRIYASEVCEDSITVGMV
RHQGKIMYVGDVRSVTQKHIQEWGPFDLVIGGSPCNDLSIVNPARKGLYEGTGRLFFEFYRLLDARPKEGDDRP
FFWLFENVVAMGVSDKRDISRFLSNPVMIDAKEVSAAHRARYFWGNLPGMNRPLASTVNDKLELQECLEHGRIA
KFSKVRTITTRSNSIKQKQDQHFVPFMNEKEDILWCTEMERVFGFPVHYTDVSNMSRLARQRLGRSWSVPVIRH
LFAPLKEYFACV

770/6881
FIGURE 719

CTGACCGCCGGGGGGTGCCCCGGGACGTAGCGCCGCGGAGAGGAAGCGGGCAAAGGGGACCATGCGGCGCCTGAC
TCGTGCGGTGTTCTGCCAGTCTTCGGGGTGCTCTGGATCACGGTGCTGCTGTTCTTCTGGGTAACCAAGAGGAA
GTTGGAGGTGCCGACGGGACCTGAAGTGCAGACCCCTAAGGTTTGGTCCTTGTTTTTCAAGGTGGCTGGGATGAG
CCCTTGGGCGCCTCAGGTGCCTGTATCACCCACTCCTCCCTACCAAAGAGGGCATCTTCCTACAGGAGGACACCT
TGCTGTATGTCATTTCCCATGTCTCTTGCAAGAAGCTCAGTTCCATTTCAGACTCAGGTCTTTCTTCAAGTCAG
ATGCACACTGCTGGTGTATTGCACGGACCTTCCACCCACTAGCATCATCATCACCTTCCACAACGAGGCCCCGCTC
CAGCTGCTCAGGACCATCCGAGTGTATTAAACCGCACCCCTACGCATCTGATCCGGGAAATCATATTAGTGGA
TGACTTCAGCAATGACCCTGATGACTGTAAACAGCTCATCAAATTGCCCAAGGTGAAATGCTTGCGCAATAATGA
ACGGCAAGGTCTGGTCCGGTCCCGGATTCCGGGGCGTGACATCGCCCAGGGCACCCTCTGACTTTCTCGACAG
CCACTGTGAGGTGAACAGGGACTGGCTCCAGCCTCTGTTGCACAGGGTCAAAGAGGACTACACGCGGGTGGTGTG
CCCTGTGATCGATATCATTAACTGGACACCTTCACCTACATCGAGTCTGCCTCGGAGCTCAGAGGGGGGTTTGA
CTGGAGCCTCCACTTCCAGTGGGAGCAGCTCTCCCCAGAGCAGAAGGCTCGGCGCCTGGACCCACGGAGCCCAT
CAGGACTCCTATCATAGCTGGAGGGCTCTTCGTGATCGACAAAGCTTGTTTGTATTACCTGGGGAAATATGATAT
GGACATGGACATCTGGGGTGGGGAGAACTTTGAAATCTCCTTCCGAGTGTGGATGTGCGGGGGCAGCCTAGAGAT
CGTCCCCTGCAGCCGAGTGGGGCAGCTCTTCCGGAAGAAGCACCCCTACGTTTTCCCTGATGGAAATGCCAACAC
GTATATAAAGAACAACCAAGCGGACAGCTGAAGTGTGGATGGATGAATACAAGCAATACTATTACGCTGCCCCGCC
ATTGCCCCTGGAGAGGCCCTTCCGGGAATGTTGAGAGCAGATTGGACCTGAGGAAGAATCTGCGCTGCCAGAGCTT
CAAGTGGTACCTGGAGAATATCTACCCTGAACTCAGCATCCCCAAGGAGTCCCTCCATCCAGAAGGGCAATATCCG
ACAGAGACAGAAGTGCCTGGAATCTCAAAGGCAGAACCAAGAAACCCCAAACCTAAAGTTGAGCCCCGTGTC
CAAGGTCAAAGGCGAAGATGCAAAGTCCCAGGTATGGGCCTTACATACACCCAGCAGATCCTCCAGGAGGAGCT
GTGCCTGTCTAGTCATCACCTTGTTCCTGGCGCCCCAGTGGTTCTTGTCTTTGCAAGAATGGAGATGACCGACA
GCAATGGACCAAACTGGTTCCACATCGAGCACATAGCATCCACCTCTGCCTCGATACAGATATGTTCCGGTGA
TGGCACCGAGAACGCAAGGAAATCGTCGTCAACCCATGTGAGTCCTCACTCATGAGCCAGCACTGGGACATGGT
GAGCTCTTGGAGACCCCTGCCAGAAGCAGCAAGGGCCATGGGGTGGTGTCTCCCTGGACCAGAACAGACTGGAAA
CTGGGCAGCAAGCAGCCTGCAACCACCTCAGACATCCTGGACTGGGAGGTGGAGGCAGAGCCCCCAGGACAGGA
GCAACTGTCTCAGGGAGGACAGAGGAAAACATCACAAGCCAATGGGGCTCAAAGACAAATCCACATGTTCTCAA
GGCCGTAAAGTTCAGTCTGCGCCAGTCATTCCCTGATTGGTATCTGGAGACAGAAACCTAATGGGAAGTGTTTA
TTGTTCTTTTCTACAAAGGAAGCAGTCTCTGGAGGCCAGAAAGAAAAGCCTTCTTTTTCTACTAGGCCAGGACT
ACATTGAGAGATGAAGAATGGAGGTTGTTTCCAAAAGAAATAAAGAGAAACTTAGAAGTTGTCTCTGG

771/6881
FIGURE 720

MRRLTRRLVLPVFGVLWITVLLFFWVTKRKLEVPTGPEVQTPKVWSLFFKVAGMSPWAPQVPVSPTPPYQRGHLPTGGHLAVCHFPCLLQEAQFHLQTQVFLQVRCTLLVYCTDLPPTSIIITFHNEARSTLLRTIRSVLNRTPTHIREIILVDDFSNDPDDCKQLIKLPKVKCLRNÑERQGLVRSRIRGADIAQGTTLTFLDSHCEVNRDWLQPLLHRVKEDYTRVVCVIDIINLDTFTYIESASELRGGFDWSLHFQWEQLSPEQKARRLDPTPIRTPIIAGGLFVIDKAWFDYLGKYDMDMDIWGGENFEISFRVWMC GGSLEIVPCSRVGHVFRKKHPYVFPDGNANTYIKNTKRTAEVWMDEYKQYYAARPFALERPFGNVESRLDLRKNLRCQSFKWYLENIYPELSIPKESSIQGNIRQRQKLESQRQNNQETPNLKLSPCAKVKGEDAKSQVWAFTYTQQILQEELCLSVITLFPGAPVVLVLCKNGDDRQWTKTGSHIEHIAASHLCLDTDMFGDGTENGKEIVVNPCESSLMSQHWMVSS

772/6881
FIGURE 721

GGTGTTCACTCAACTTGGATCTGTGCTGAAAAATTGTGACATTTTCAGTACATCTGGTAGAGGGTACAGCTTTTAT
CTTGACATGAATTTTTTGTGTTCTTCCTGTGCATAAATTCAGAAAACACCACAGGGATGGCGAGAAGTATTT
GTTGACATTGATCCACAGGTTTCTGATAAACTGAGGTTTGTTTTGGCACCTTCTGCCACCCACAGCAGAAGCCTTC
ATACAACATGACGAAACAAGGGATCATGTTGAAGTGTGTCTGATGCTGGTGTATCATCGAGGAACCTTTCTCAA
CGCATTGCATTAACTGGAGGTGCTGCACTGGTTGCTGATTATGGTCATGATGGAACAAAGACAGATACCTTCAGA
GGGTTTTGCGACCACAAGCTTCATGATGTCTTAATTGCCCCAGGAACAGCAGATCTAACAGCTGATGTGGACTTC
AGTTATTTGCGAAGAATGGCACAGGGAAAAGTAGCCTCTCTGGGCCCAATAAAACAACACACATTTTTAAAAAAT
ATGGGTATTGATGTCCGGCTGAAGGTTCTTTTAGATAAATCAAATGAGCCATCAGTGAGGCAGCAGTTACTTCAA
GGATATGATATGTTAATGAATCCAAAGAAGATGGGAGAGAGATTTAACTTTTTGCCTTGCTACCTCATCAGAGA
CTTCAAGGTGGAAGATATCAGAGGAATGCACGTCAGTCAAACCCTTTGCATCCGTTGTAGCTGGGTTTAGTGAA
CTTGCTTGGCAGTGATATTTTCAGCTTGGACATTTTACCCTTCAGTCGGCCCAAGAAATCAAATAAAGGAAACAC
ATTTTCATATACTGCAGGTAACAAAAGTCAAAGTATTTTATCTTTTCACAGCAAGAACAGTCCATGTTGTATATAA
TACAACCAACATTATAGAAGCTTTAGGGTTGTGACTGGCTTTGGTGCAAATGTGTGCTCAAGCTAATAAGTTATT
GTGAAACTGAGTTTCCTTTAACTTACAAAGCTAGTTGCCATATTTCTATTTTATTTTAAAAAGTAAACATGCGGC
TGGGCGTGGTGGCTCATGCCTGTAATCCCAGCACTTTGGGAGGCTGAGGTGGGCATATCACCTGAGGTCAGCAGT
TAAAGACCAGCCTGACCAAAATGGAGAAACCCCATCTCTACTAAAAATACAAAACCTAGCCGGGTATGGTGGTACA
TGCTGTAAATCCCAGC

773/6881
FIGURE 722

MAQGKVASLGPIKQHTFLKNMGIDVRLKVLLDKSNEPSVRQQLLQGYDMLMNPKKMGERFNFFALLPHQRLQGGR
YQRNARQSKPFASVVAGFSELAWQ

774/6881
FIGURE 723

CTTCTCTCGCCAGGCATCCTCGTGGAAGTGACATCGTCTTTAAACCCTGCGTGGCAATCCCTGACGCACCGCCGT
GATGCCCCAGGGAAGACAGGGCGACCTGGAAGTCCAACACTACTTCCTTAAGATCATCCAACACTATTGGATGATTATCC
AAAATGTTTCATCGTGGGAGCAGACAATGTGGGCTCCAAGCAGATGCAGCAGATCCGCATGTCCCTTCGCGGGAA
GGTCGTGGTGCTGATGGGCAAGAACACCATGATGCGCAAGGCCATCCGAGGGCACCTGGAAAACAACCCAGCTCT
GGAGAACTGCTGCCTCATATCTGGGGGAATGTGGGCTTTGTGTTACCAAGGAGGACCTCACTGAGATCAGGGA
CATGTTGCTGGCCAATAAGGTGCCAGCTGCTGCCCCTGCTGGTGCCATTGCCCCATGTGAAGTCACTGTGCCAGC
CCAGAACACTGGTCTCGGGCCCGAGAAGACCTCCTTTTTCCAGGCTTTAGGTATCACCACTAAAATCTCCAGGGG
CACCATTGAAATCCTGAGTGATGTGCAGCTGATCAAGACTGGAGACAAAGTGGGAGCCAGCGAAGCCACGCTGCT
GAACATGCTCAACATCTCCCCCTTCTCCTTTGGGCTGGTTCATCCAGCAGGTATTCGACAATGGCAGCATCTACAA
CCCTGAAGTGCTTGATAAAACAGAGGAACTCTGCATTCTCGCTTCCTGGAGGGTGTCGCAATGTTGCCAGTGT
CTGTCTGCAGACTGGCTACCCAACTGTTGCATCAGTACCCCATCTATCATCAACGGGTACAAACGAGTCCTGGC
CTTGCTGTGGAGACGGATTACACCTTCCCCTTCTGCTGAAAATGTCAAGGCCTTCTTGGCTGATCCATCTGCCTT
TGTGGCTGCTGCCCCCTGTGGCTGCTGACACCACAGCTGCTCCTGCTGCTGCTGCAGCCCCAGCTAAGGTTGAAGC
CAAGGAAGAGTCGGAGGAGTCGGACGAGGATATGGGATTTGGTCTCTTTGACTTAATCACCAAAAAGCAACCAACT
TAGCCAGCTTTATTTGCAAAACAAGGAAATAAAGGCTTACTTCTTT

775/6881
FIGURE 724

MPREDRATWKSNYFLKIIQLDDYPKCFIVGADNVGSKQMQQIRMSLRGKVVVL MGKNTMMRKAIRGHLENNPAL
EKLLPHIWGNVGFVFTKEDLTEIRDMLLANKVPAAARAGAIAPCEVTVPAQNTGLGPEKTSFFQALGITT KISRG
TIEILSDVQLIKTGDKVGASEATLLNMLNISPF SFGLVIQQVFDNGSIYNPEVLDKTEETLHSRFL EGVRNVASV
CLQTYPTVASVPHSIINGYKRVLALSVETDYTFPLAENVKAFLADPSAFVAAAPVAADTTAAPAAAAAPAKVEA
KEESESEDEDMGFGLFD

776/6881
FIGURE 725

GTCTTTAAACCCTGCGTGGCAATCCCTGACGCACCGCGTGATGCCCAGGGAAGACAGGGCGACCTGGAAGTCCA
ACTACTTCCTTAAGATCATCCAACCTATTGGATGATTATCCAAAATGTTTCATCGTGGGAGCAGACAATGTGGGCT
CCAAGCAGATGCAGCAGATCCGCATGTCCCTTCGCGGGAAGGTCGTGGTGCTGATGGGCAAGAACACCATGATGC
GCAAGGCCATCCGAGGGCACCTGGAAAACAACCCAGCTCTGGAGAACTGCTGCCTCATATCTGGGGGAATGTGG
GCTTTGTGTTACCAAGGAGGACCTCACTGAGATCAGGGACATGTTGCTGGCCAATAAGGTGCCAGCTGCTGCCC
GTGCTGGTGCCATTGCCCCATGTGAAGTCACTGTGCCAGCCCAGAACACTGGTCTCGGGCCCGAGAAGACCTCCT
TTTTCCAGGCTTTAGAGTGATGTGCAGCTGATCAAGACTGGAGACAAAGTGGGAGCCAGCGAAGCCACGCTGCTG
AACATGCTCAACATCTCCCCCTTCCTTTGGGCTGGTCATCCAGCAGGTATTGACAAATGGCAGCATCTACAAC
CCTGAAGTGCTTGATAAAACAGAGGAACTCTGCATTCTCGCTTCCTGGAGGGTGTCCGCAATGTTGCCAGTGTC
TGTCTGCAGACTGGCTACCCAACCTGTTGCATCAGTACCC

777/6881
FIGURE 726

CTGCGTGGCAATCCCTGACGCACCGCCGTGATGCCCAGGGAAGACAGGGCGACCTGGAAGTCCAACACTTTCCTT
AAGATCATCCAACCTATTGGATGATTATCCAAAATGTTTCATCGTGGGAGCAGACAATGTGGGCTCCAAGCAGATG
CAGCAGATCCGCATGTCCCTTCGCGGGAAGGTCGTGGTGCTGATGGGCAAGAACACCATGATGCGCAAGGCCATC
CGAGGGCACCTGGAAAAACAACCCAGCTCTGGAGAACTGCTGCCTCATATCTGGGGGAATGTGGGCTTTGTGTTC
ACCAAGGAGGACCTCACTGAGATCAGGGACATGTTGCTGGCCAATAAGGTGCCAGCTGCTGCCGTGCTGGTGCC
ATTGCCCCATGTGAAGTCACTGTGCCAGCCCAGAACACTGGTCTCGGGCCCGAGAAGACCTCCTTTTTCCAGGCT
TTAGGTATCACCCTAAAAATCTCCAGGGGCACCATTTGAAATCCTGACAGGTATTGACAATGGCAGCATCTACAA
CCCTGAAGTGCTTGATAAAACAGAGGAACTCTGCATTCTCGCTTCCTGGAGGGTGTCCGCAATGTTGCCAGTGT
CTGTCTGCAGACTGGCTACCCAACCTGTTGCATCAGTACCCCATTTCTATCATCAACGGGTACAAACGAGTCCTGGC

778/6881
FIGURE 727

CCTCTAGCTTAGCGAGCGCTGGAGTTTGAAGAGCGGGCAGTGGCTGCACACGCCAAACTTTCCCTATGGCTTCGG
TGACCAGGGCCGTGTTTGGAGAGCTGCCCTCGGGAGGAGGGACAGTGGAGAAGTTCCAGCTGCAGTCAGACCTCT
TGAGAGTGGACATCATCTCCTGGGGCTGCACGATCACAGCCCTAGAGGTCAAAGACAGGCAGGGGAGAGCCTCGG
ACGTGGTGCTTGGCTTCGCCGAGTTGGAAGGATACCTCCAAAAGCAGCCATACTTTGGAGCAGTTATTGGGAGGG
TGGCCAACCGAATCGCCAAAGGAACCTTCAAGGTGGATGGGAAGGAGTATCACCTGGCCATTAACAAGGAACCCA
ACAGTCTGCATGGAGGAGTCAGAGGGTTTGATAAAGTGCTCTGGACCCCTCGGGTGCTGTCAAATGGCGTCCAGT
TCTCGCGCATCAGTCCAGATGGTGAAGAAGGCTACCCCGGAGAGTTAAAAGTCTGGGTGACATACACCCTGGATG
GCGGAGAGCTCATAGTCAACTACAGAGCACAAGCCAGTCAGGCCACACCAGTCAACCTGACCAACCATTCTTACT
TCAACCTGGCAGGCCAGGCTTCCCAATATAAATGACCATGAAGTCACCATAGAAGCGGATACTTATTTGCCTG
TGGATGAAACCCTGATTCTACAGGAGAAGTTGCCCCAGTGAAGGCACTGCATTTCGACCTGAGAAAGCCAGTGG
AGCTTGGAAAACACCTGCAGGACTTCCATCTCAATGGTTTTGACCACAATTTCTGTCTGAAGGGATCTAAAGAAA
AGCATTTTTGTGAAGGGTGCATCATGCTGCAAGCGGGCGGGTACTAGAAGTATACACCACCCAGCCCGGGTCC
AGTTTTACACGGGCAACTTCCTGGATGGCACATTAAAGGGCAAGAATGGAGCTGTCTATCCCAAGCACTCCGGTT
TCTGCCTGGAGACTCAGAACTGGCCTGATGCAGTCAATCAGCCCCGCTTCCCTCCTGTGCTGTCTGAGGCCTGGTG
AGGAGTATGACCACACCACCTGGTTCAAGTTTTCTGTGGCTTAAGGAAGTGTGAAGATATGATCCAGTCCAGGGC
TAGGCTCAGCCACCTGTCTCCTGTCCAGAAAAAGGTGAAGATTAAGAAGCTTTCAGAAATGATTCTATGGATTAA
AATCATACAAATGGTGGCTGTTCTGAGAATCAGTCTGGGTATTGATTTCTTTTCCAGTGAAGTCCAGGCCA
TGTCTAATGACCAGCTCGATTCCCTGTGCAGTTCAGAGAGCAAGTGAACCAACCAACAATGTCGTCTATCTAAGC
CTTGACCCTAGCCAGGGACTCCCATGTCTGTGTTGGCTCCATCTCTCCACACTGCCTCTTTCTTTTCAACTTTTT
GCCCTTCCTTTCTTTAAAGCTATTCTCACATTGCTTTTATTTTCTCCTCCTTCACCTCCAACCACTGTCTAGCAGC
ACTCTGGAGTTTTCAAATGTACATTAGCCTCACCCTGCATGCTAGGAGATGGACCTGTCTCTATACAGCAGTAG
ATGATTGATAAGTGAGGAACTGAGGCTTACAGAGGTTGAAGACCAACAAGCTAATAAATAAAAAGTTGAGGCCG
GGCAGGCTGGCTCACGCCTGTAATCCCAGCACTTTGGAAGGCCGAGGCAGGCGGATCACGAGGTCAGGAGATCGA
GACCATCCTGGCTAACACGGTGAAACCCCGTCTCTACATAAAAAATTAGCTGGGTGTGGTAGCACGCACCCGTAG
TTCCAGCTACTTGGGAGGCTGAAGCAGGAGAATCGCTTGAACCTGGGAGGTGGAGGTTGCAGTGAGCCGAGATCA
TGCCACTGCAGTGGGCAACAGAGCAAACTCGATCTCAAAAAAAAAAAAAAAAAAAAAA

779/6881
FIGURE 728

MASVIRAVFGELPSGGGTVEKFQLQSDLLRVDIISWGCTITALEVKDRQGRASDVVLGFAELEGYLQKQPYFGAV
IGRVANRIAKGTFKVDGKEYHLAINKEPNSLHGGVVRGFDKVLWTPRVLSNGVQFSRISPDGEEGYPGELKVWVTY
TLDGGELIVNYRAQASQATPVNLTNHSYFNLAGQASPNINDHEVTIEADTYLPVDETLIPTGEVAPVQGTAFDLR
KPVELGKHLQDFHLNGFDHNFCLKGSKEKHFCARVHHAASGRVLEVYTTQPGVQFYTGNFLDGTLKGKNGAVYPK
HSGFCLETQNWPDVAVNQPRFPPVLLRPGEEDHTTWFKFSVA

780/6881
FIGURE 729

GACTTCAGGAAAAATCAACCCAGCACCAGCAGCTACCAACCACCATTCCATCTCTTCACTTGAACAGCATTAGTT
AAGTCCAGATGTGGGAACCCCTTCTCTTGGAAGAAGTTCCTAATTGTGTCTCAGACCGGTGTAAACAAACCAGCCA
GCCGCCACCTTGCTAAACCTATAAGCTTTTTTAAATCCAATATATTCTGCCAAGAATATGCCTTGATAGTTAGCC
CTCAGCCCATAGGTGTTTTTTGTTTTTAAACAGAATTATATATGTCTGGGGGTGAAAAACCCCTTGCATTCCAAA
GGTCCATACTGGTTACTTGGTTTTATTGCCACCACTTAGTGGATGTTTCACTTTAGAACCATTTTGTCTGCTCCCT
CTGGAAGCCCTTGCGCAGAGCTTACTTTTGTAAATTGTTGGAGAATAACTGCTGAATTTTAGCTGTTTTGAGTTGAT
TCGCACCACTGCACCACAACCTCAATATGAAAACCTATTTAACTTATTTATTATCTTGTGAAAAGTATACAATGAAA
ATTTTGTTCATACTGTATTTATCAAGTATGATGAAAAGCAATAGATATATATCTTTTATTATGTTAAATTATGA
TTGCCATTATTAATCGGCAAAATGTGGAGTGTATGTTCTTTTACAGTAATATATGCCTTTTGTAACTTCACITG
GTTATTTTATTGTAAATGAGTACAAAATTCTTAATTTAAGAGATTGTATGTAATATTTATTTTCAATATTTCTTT
CCTTGTTTACGTAAATTTTGAAAGATTGCATGATTTCTTGACAGAAATCGATCTTGATGCTGTGGAAGTAGTTG
AGGAACATCCTATGAGTTTTCTTAGAATGTATAAAGGTTGTAGCCCATCCAACCTCAAAGAAAAAATGACCACA
TACTTTGCAATCAGGCTGAAATGTGGCATGCTTTTCTAATTCCAACCTTTATAAACTAGCAAAAAAGTGTGCTT
ATTCCACCAGTTCTACTGTGACATACTCGAGTATAAAGACATGTAGCAATAACGGGGAGTGGGGGGGAGTCTCA
CAGTGCCTTTGGAAGGGCCCGAACTTGCCTTAAATCTTCCTCAACCAATAAGTATTTTATTAGTGCTTGAGAGA
ATCTGAATGTAGGATGGGTTCAACTGCACAAAAGGAAAAGATTTTACCACCTTTTTTATATAGATATAAAGTGA
AGCAACCGCCTTAGTGCTGAAATATGTAGTACATGAATATGCCTTGTTTAAATTACAGAAAATTCCTTCCTGGACA
GCAGATGAATGAGCGGTAGCTTTAGTTTGTACGTAGGTACAGTTGGAGCACTATATGTACTCTCTGGACTACTTT
GGACAGAAGTAGGTTTTTGAATGTAACAAGATAAGTCAACTTGAGTTGTAATATATTTTGGGGAATCAGCTCACT
ACAAATTGTGACTGTAAACATTGTACTGTAAATGTTTTGTAGTT

781/6881

FIGURE 730

MWEPFSWKKFLIVSQTGVNKPASRHLAKPISFLKSNIFCQEYALIVSPQPIGVFCFLTELYMSGGEKTLAFQRSI
LVTWFHCHHLVDVQFRTILSAPSGSLAQSLCNCWRITAEFLAVLS

782/6881
FIGURE 731

CCTGATTCTAGGCTGGTCACTACTCCGAGCCTGTGACGTTTGC GGCAGCCAGGCCGTCGACGATGCCAGTGAAA
CTCTCTGGGAAATTGCAAAAGCTGAAGTGGAAAAAAGGGGAATTAATGGAAGTGAAGGTGATGGAGCTGAAATTG
CAGAAAAATTTGTTTTCTTCATTGGCAGTAAAAATGGGGGAAAGACTACTATTATTCTAAGGTGTCTTGACAGAG
ATGAACCACCAAAACCAACCTTAGCTTTGGAATATACATATGGAAGAAGAGCAAAAGGGCACAACACACCAAAAG
ATATCGCTCACTTTTGGGAACCTCGGTGGAGGAACCTCTTTATTGGACTTAATCAGCATACCCATCACAGGTGACA
CCTTACGGACGTTTTCTCTTGTTCTCGTTCTGGATCTTTCAAAACCTAATGATCTCTGGCCCACCATGGAAAATC
TCTTGCAAGCCACAAAAGCCATGTAGACAAAGTGATAATGAAACTGGGAAAGACAAATGCTAAAGCAGTTTCTG
AAATGAGACAGAAGATCTGGAATAATATGCCGAAGGATCATCTGATCATGAATTAATTGACCCATTTCCGGTAC
CTCTGGTCATAATTGGAAGTAAATATGATGTTTTTTCAGGATTTTGAGTCTGAGAAGAGAAAGGTAATATGCAAGA
CACTTCGATTTGTTGCACATTATTATGGAGCATCATTAAATGTTTACCAGTAAATCAGAAGCTCTATTACTAAAAA
TACGTGGAGTTATCAACCAGTTGGCATTGGCATTGACAAAAGCAAATCAATATGTGTGGATCAGAATAAACCGC
TGTTTATCACAGCAGGATTGGATTCTTTCGGTCAAATAGGATCTCCTCCTGTTCTGAAAATGACATTGGAAAGC
TTCATGCCCCACTCACCTATGGAGTTGTGGAAAAAAGTGATGAAAAGCTCTTCCACCAAGAGTATTAACACGC
TGAAAGATATCAAGGACCTGCGAGAGATCCTCAGTATGCTGAAAATGAAGTCGATGAGATGAGAATTCAGAAGG
ATCTGGAACCTGGAACAGTACAAAAGAAGTTCTTCCAAGTCTTGGAACAAATCGAGCTTGATTCTTGAACCTATT
TCAATTATTGTATATTTATTCTTTCTTTTCCAAATACAAATAAGATTATACTGTGAATTAATCTATTGTGGCAATA
TGTGAAGAAAGTTAAACTGTATAATTTGTTAAAGGACAAGCTGGATTTCTTGGACTAGTGCATCTCCCTGTATAT
CTTGAAGCTTTTTTAAAAGGAAAAATTATTGTAGAACCACGTGTAATTTTTTTTTTAAAATAAAAGAAATCTTCTACTA
CCTACCTCT

783/6881
FIGURE 732

MPSETLWEIAKAEVEKRGINGSEGDGAEIAEKVFVFFIGSKNGGKTTIILRCLDRDEPPKPTLALEYTYGRRAKGH
NTPKDIAHFWEELGGGTSLLDLISIPITGDTLRTFSLVLVLDLSKPNDLWPTMENLLQATKSHVDKVIMKLGKTNA
KAVSEMRQKIWNMPKDHDPHELIDPFPVPLVIIGSKYDVFQDFESEKRKVICKTLRFVAHYYGASLMFTSKSEA
LLLKIRGVINQLAFGIDKSKSICVDQNKPLFITAGLDSFGQIGSPVPENDIGKLHAHSPMELWKKVYEKLFPPK
SINTLKDIKDPARDPQYAENEVDEMRIQKDLELEQYKRSSSKSWKQIELDS

784/6881
FIGURE 733A

ACATGCTCCTCCTGTCTTCTGGCGGAGCGTGCTTCCCGCTGCGGGGACGTTTCGAGCAATGGCAGCCCTGCTGAG
ATCCGCGCGTTGGTTGCTGCGTGCCGGGGCGGCCCGCGCCTCCCGCTCTCCCTGCGCCTCCTCCCTGGCGGGCC
GGGCGCGCTGCATGCCGCTCCTATCTGCCC GCCGCTCGCGCCGGGCCCGTGCCGGAGGACTACTGAGCCCAGC
CAGGCTGTATGCCATTGCTGCCAAAGAAAAAGATATTCAAGAGGAGTCCACTTTTTCTTCTAGGAAGATTTCCAA
TCAGTTTGATTGGCTCTAATGAGACTAGATCTTTCTGTTTCAAGAACTGGCCGCATTCCAAAGAAGCTTCTACA
AAAAGTTTTTAATGATACCTGCCGCTCAGGTGGCCTAGGTGGTAGTCATGCCCTTGCTTCTACTACGTAGTTGTGG
TTCTCTCTTGCCCTGAAGCTTGAAGAGAGAACAGAATTTGCTCATAGGATATGGGACACACTTCAGAAATT
AGGTGCTGTGTATGATGTGAGTCACTATAATGCTTTACTTAAAGTCTATCTTCAAAATGAATATAAAATTCTCACC
AACTGATTTTCTGGCAAAAATGGAGGAAGCAAAACATTCAACCAATCGAGTGACATACCAGAGATTGATTGCTTC
TTATTGTAATGTAGGAGATATTGAAGGTGCCAGCAAGATTCTTGGATTATGAAAATAAGGATCTCCAGTTAC
AGAGGCAGTATTCAGTGCCCTTGTGACAGGGCATGCCAGAGCTGGTGATATGGAGAATGCAGAAAACATTCTCAC
AGTGATGAGAGATGCCGGAATTGAGCCTGGTCCAGACACATACCTCGCATTATTGAATGCATATGCTGAGAAGGG
CGACATTGACCATGTTAAGCAGACTCTGGAGAAGGTGGAGAAGTCCGAGCTTCACCTTATGGACCGTGATTTACT
GCAAATTATTTTTAGCTTCAGTAAAGCTGGGTATCCTCAGTATGTCTCAGAAAATTTGGAAAAAGTTACATGTGA
AAGAAGATATATTCCAGATGCAATGAACCTCATTTTTACTTTTAGTCACTGAAAAATTTGGAAGATGTAGCGTTGCA
AATTTTACTAGCATGCCCCGTATCAAAGGAAGATGGCCCAAGTGTCTTTGGCAGTTTCTTTTTACAACACGTGTGT
GACTATGAATACGCCGTGTGGAGAAGCTAACAGACTACTGTAAGAAGTTAAAGGAAGTCCAGATGCACTCCTTTCC
TCTGCAGTTCACCTCCATTGTGCTTTACTCGCCAATAAACTGATTTGGCAAAAAGCCTTAATGAAGGCTGTGAA
GGAGGAAGGTTTTCTATCAGACCTCACTATTTCTGGCCATTGCTAGTTGGACGTCGGAAGGAAAAAATGTTCA
AGGTATAATTGAAATCCTCAAAGGAATGCAAGAATTGGGAGTACATCCTGATCAGGAAACATATACAGATTATGT
GATTCCATGCTTTGATAGTGTAAGTCAAGCAGAGCCATTTTGCAAGAAAATGGATGTCTGTCTGATAGTGATAT
GTTTTCTCAAGCTGGATTGAGAAGTGAAGCAGCAAAATGGGAACCTTAGACTTTGTATTATCATTTTTGAAATCAA
TACATTGCCCATCTCGCTGCAGTCTATAAGAAGTAGCCTACTGCTAGGCTTCAGGAGGTCTATGAATATAAATCT
TTGGAGCGAGATAACAGAATTGTTGTACAAGGATGGACGTTATTGCCAGGAGCCTCGAGGACCGACGGAAGCTGT
TGGCTATTTTTCTTTATAACTTGATTGACAGCATGAGTGACTCAGAGGTACAGGCCAAGGAGGACATTGTAGACA
ATACTTCCATCAGCTGGAGAAGATGAATGTAAAAATTCCTGAAAATATCTACAGAGGCATTCGTAATCTCCTGGA
AAGCTACCATGTTCCGAAATTGATTAAAGGATGCTCACTTGTGGTTGAGAGTAAGAATTTAGACTTTCAAAAAAC
TGTGCAACTTACATCATCTGAATTGGAGTCCACACTTGAACACTAAAAGCTGAAAATCAACCTATAAGAGATGT
CCTAAAGCAACTCATATTAGTGCTTTGTTTCAAGAGAATATGCAAAAAGCCCTGAATTGAAAGCAAAATATGA
ATCCGACATGGTTACTGGTGGCTATGCAGCTTTAATAAATTTATGCTGTGACATGATAAAGTAGAAGATGCCTT
GAACTTGAAAGAAGAAATTTGACCGCTTAGATTCACTGCTGTGCTTACACCGCAAGTATGTAGGCCTTGTAAG
AGTATTGGCAAAGCATGGCAAGCTCCAAGATGCTATTAACTTCTGAAGGAGATGAAAGAGAAGGATGTTCTTAT
CAAAGATACAACAGCCTTGTCTTTTTCCACATGCTAAATGGCGCAGCTTTAAGAGGTGAAATTGAAACAGTAAA
ACAGTTGCATGAAGCCATCGTGACTCTAGGGTTAGCAGAACCATCCACCAACATAAGTTTCCCATTTGGTCACTGT
ACACTTGGAAAAGGGCGACCTATCTACTGCTCTTGAGGTCGCCATTGACTGCTATGAAAAGTATAAAGTATTACC
AAGGATTCATGATGCTTGTGTAACTGGTAGAGAAAGGCGAGACTGATCTAATTCAGAAAGCAATGGACTTTGT
GAGCCAAGAACAAGGTGAAATGGTGATGCTCTATGATCTCTTCTTTGCCTTCTTACAAACAGGAAATTACAAAGA
GGCCAAGAAGATCATTGAGACTCCAGGGATTAGAGCTCGATCTGCAAGGCTTCAGTGGTTTTGTGACAGATGTGT
TGCAATAATCAGGTTGAACTCTGAAAAATTAGTGGAGCTGACACAGAAGCTATTGAAATGTGATAGAGACCA
GATGTACTACAATCTGTAAAATGTATAAAATAAACGGTGACTGGCAAAGAGCTGATGCAGCTGGAATAAAAT
CCAAGAAGAAAATGTTATCTCTCGTGAAAAGACATTAAAGATTATTAGCAGAAATCCTTAGAGAGGGTAAACCAGGA
AGTTCCGTTTGACGTACCTGAGTTGTGGTATGAAGATGAAAAACATTCCCTGAATTCTTCGTCAGCCTCAACCAC
AGAACCTGATTTCCAGAAAGATATATTGATTGCCTGCCGATTGAACCAAAAAAAGGGGCATATGATATTTTCT
GAATGCAAAAGAGCAAAACATTGTGTTTTAATGCTGAAACCTACAGCAATCTCATTAAATTACTGATGTGAGAAGA
TTATTTTACACAAGCAATGGAAGTGAAGCATTGCGGGAGACCCACATCAAGGGCTTCACACTGAACGATGCTGC
CAACAGCCGCTCATCATAACGCAAGTTAGCGGGGATTATTTGAAAGAGGCTGTGACAACACTGAAACAGTATT
GGATCAGCAGCAGACCCCTTCTAGGTTAGCAGTGACCGGTGTCATCCAGGCATTGGCCATGAAGGGTGATGTTGA
AAACATAGAAGTAGTTTCAAGATGTTAAATGGACTCGAAGACTCCATTGGACTTTCAAAAATGGTTTTTCATCAA

785/6881

FIGURE 733B

TAACATTGCTTTGGCTCAAATAAAGAATAATAACATAGATGCCGCAATAGAAAACATTGAAAATATGCTTACTTC
AGAGAATAAAGTCATTGAACCCCAATACTTCGGCTTGGCATACTTATTCAGAAAAGTAATAGAGGAGCAGTTGGA
ACCAGCAGTTGAAAAGAGTAACATCATGGCGGAGAGATTGGCCAATCAGTTTGCAATTTATAAACCTGTCACTGA
TTTTTTCCTTCAACTTGTGGATGCAGGCAAGGTGGATGATGCCAGAGCTCTCCTACAGAGATGTGGTGCAATTGC
TGAACAAACCCCGATTTTGTGTTGTTTCCTCCTTAGGAATTCTAGGAAACAAGGAAAGGCATCAACTGTGAAATC
TGTGTTAGAATTGATTCCCTGAATTAAATGAAAAGGAAGAAGCATACAATTCCCTCATGAAAAGCTATGTCTCAGA
GAAAGATGTACATCTGCTAAAGCACTGTATGAACATTTGACTGCAAAGAATACAAAATTGGATGATCTGTTTCT
AAAGCGTTACGCATCTTTGCTGAAGTATGCTGGAGAGCCTGTCCCTTTCATTGAACCCCTGAAAGCTTTGAATT
TTATGCACAGCAGCTAAGAAAATTGAGGGGAAAACCTTCTTGAATAACCAGGCGATACTTTGTTTTGTATATAT
TTGTGATTCTGTGTCTACAIGTTATTTTGAAGTATATCTGAGGGGAAAATAAATGAAAATTTTCTTTATGTACTT
ATGTATGTGTGATGCATGTTCAAAGTCTTATTGACCATAACTCTGTGCACTTGGTTATTGGACATTTTGGAGTT
TTTTCTCTGGGAAAAATCGATAGTGTTTTCTTCAATGCTGCTGTGTGTAAGCCATACTTTTTTCAGGATTCTT
CCCTAATTGGCTCTTTGGTTTTCCCTGCTCTGTTTCATTTATTTTATTAAATGTTATTCTTTATTTAAGATTCT
ACTTATTAGTCTGCTGTTTCTCTGAAAAATTTTAGAGCTAGGTATAGTGACCGTGAACCTTTCTAACGCATAATAT
TCTGTGATACAGCCATTCCGTACATGTGTGAAGTCCTGCATAACTTTTGAACCTTTGTTAAATGTTGGCACTAGGA
GTCATCAGATCTAGGCTTCATCATTTTCCAGTGAGAAGCAGAGACCCAAAGGGCCTGTTACTTGTGCTTGGTCAG
GGGACTGTCTGTATGCCTGGAGGCTCTTCGGCACACTTCCCATCTTTCCCTTCTGCCACTGTGGCTTCAAGCA
CCTCTGTTTCATAGAGCGTCTCTGAAATTGAGTCTCGGTCATGACTTATCCCGAAGTAGAGCAATGTGTTTCTCT
CATTGTAGTTTCAAGGACTTTGTGAGTACAAGCTCTGCCCTAGGCTTGTTACTTTTATACTCATATCCTGAAAAGAT
GTGATTTCATCTATGAAGGGGTAAAATATTGGTTTGTATTTAATTGTTTGAATAAAAAGTGATCCCTATATTG

786/6881
FIGURE 734

ATGGGCTCAGGCTACACCAAGGCCGTGTACCGGGTCCGCCTGCCCGGCGGTGCCGCGGTGGCGCTCAAGGCGGTG
GACTTTAGCGGCCACGATCTGGGCAGCTGCGTGCGCGAGTTCCGGGTACGGAGGGGCTGCTATCGGCTGGCGGCC
CACAAAGCTGCTTAAGGAGATGGTGCTGCTGGAGCGGCTGCGGCACCCCAACGTGCTGCAGCTCTATGGCTACTGC
TACCAGGACAGCGAGGACATCCCAGACACCCTGACCACCATCACGGAGCTGGGCGCCCTGTAGAAATGATCCAG
CTGCTGCAAACTTCCTGGGAGGATCGATTCCGAATCTGCCTGAGCCTGGGCGCCCTCCTCCACCACCTGGCCAC
TCCCCACTGGGCTCCGTCACCTCTGCTGGACTTCCGCCCTCGGCAGTTTGTGCTGGTGGATGGGAGCTCAAAGTG
ACGGACCTGGATGACGCACGTGTGGAGGAGACGCCGTGTGCAGGCAGCACCGACTGCATACTCGAGTTTCCGGCC
AGGAACTTCACCTGCCCTGCTCAGCCCAGGGCTGGTGCGAGGGCATGAACGAGAAGCGGAACCTCTATAATGCC
TACAGGTTTTTCTTCACATACCTCCTGCCTCAGAGTCCCCGCCTTCACTGCGTCCTCTGCTGGACAGCATCGTC
AACGCCACAGGAGAGCTCGCCTGGGGGGTGGACGAGACCCTGGCCCAGCTGGAGAAGGTGCTGCACCTGTACCGG
AGCGGGCAGTATCTGCAGAACTCCACGGCAAGCAGCAGTACCGAGTACCAGTGTATCCCAGACAGCACCATCCCC
CAGGAAGACTACCGCTGCTGGCCATCCTACCACCACGGGAGCTGCCTCCTTTTCACTGTTCAACCTGGCTGAGGCT
GTGGATGTCTGTGAGAGCCATGCCCAGTGTGGGCTTTGTGGTCACCAACCAGACCACCTGGACAGGTGCGGCAG
CTGGTCTTTTTCAAGACTGGATGGAGCCAAGTGGTCCCTGATCCCAACAAGACCACATATGTGAAGGCCTCTGGC
TGA

787/6881
FIGURE 735

MSGYTKAVYRVRLPGGAAVALKAVDFSGHDLGSCVREFGVRRGCYRLAAHKLLKEMVLLERLRHPNVLQLYGYC
YQDSEDIPDILTTITELGAPVEMIQLLQTSWEDRFRICLSLGRLLHHLAHSPLGSVTLLDFRPRQFVLVDGELKV
TDLDDARVEETPCAGSTDCILEFPARNFTLPCSAQGWCEGMNEKRNLYNAYRFFFTYLLPHSAPPSLRPLLDIV
NATGELAWGVDETLAQLEKVLHLYRSGQYLQNSTASSSTEYQCIPDSTIPQEDYRCWPSYHHGSCLLSVFNLAEA
VDVCESHAQCRAFVVTNQTWTGRQLVFFKTGWSQVVPDPNKTTYVKASG

788/6881
FIGURE 736

GCTAACATCCCAGACAGACAGATGTGACCAGGACAAACGTGCAATAATGCCAAATGTTAAAATGTGAGTTTACCA
GCCTAGCTATGGGACTGCTGGCTCCTAGTCCAGGAATCATGGGGGTATGACTGCCTCTCCAACCCTGTGGGCTGT
AAGCAAGCTCAGGCTAGTCTCCCCACTGGGGGCTGTGCCCTCCCTGGGACGGTTCGTTGGGCAGCCCCATCACT
GTGTTCAATAGTGTGAGAATGTAGCTGCGTGGGGACAATCCATCGTGGAGTGTCTCTCAGCTTAGGTCTGGACA
GGAGACTTGGCGGGAGATGCTCCAGGATGTGGGTGATTCTGTACCTGGGGAGGCTATCTCTGACCTCCCGACAGG
GGACACTCCCAGGCCAGCCAGGGGTGAGGGGCAGAGGTGCACACCTCAGCATGAGCCAAGACTGGGGTCAGGGA
GCAGGTGTGGTTTGAGCCAGGACCTGGGGCGGGGGTGGGGCCGGGGCCTTTCTGCCTCATTGTCTTCAATGAAA
GCCTCAAAGCAGCCAAAACCAGGCTTTCCCCCTTCCTCGAGTTTGAATATCCAGAATCTTTGTACTTCTTGTG
GTTAAATTGTTTATTTTTGTAAAAATAAAATAAAATTAGTTAATAA

789/6881
FIGURE 737

MWVILYLGRSLTSRQGILPGQPRGQGQRCTPQHEPRLGSGSRCGLSQDLGRGWGRGLSASFANESLKAAKTRL
SPFLEFEYPESFVLLVG

790/6881
FIGURE 738A

AGATGACTTTGGGAAATTGCTGCTGGCTGAGGCCCTCCTGGAGCAGTGTGTTGAAGGAGAACCATGCCAAAATAAA
AGACTCCATGCCTTTGCTGGAGAAGAATGAGCCGAAGATGAGCGAAGCCAAAATTATCTAAGCAGTATCCTTAA
CCATGGGAGGCTCTCGCCACAGTACATGTGTGAGGCCATGCTGATCCTGGGCAAACCTGCATTACGTGGAGGGCTC
ATACCGAGATGCCATCAGCATGTACGCACGGGCCGGGATTGATGACATGTCCATGGAGAACAAGCCCCGTATCA
GATGCGGCTGCTGTCGGAGGCTTTTGTCACTCAAAGGCCTCTCTCTGGAACGCCTACCCAACTCCATCGCCTCCCG
CTTCCGCCTGACAGAGAGGGAGGAGGAAGTGATCACCTGTTTTGAGAGGGCCTCCTGGATCGCTCAGGTGTTCT
GCAGGAATTGGAGAAGACCACAAATAACAGCACGTCGAGGCATCTGAAAGGCTGTACCCGCTTGACTATGAGCT
CACCTACTTCTGGAAGCTGCCCTCCAGAGCGCCTATGTGAAAACCTGAAGAAGGGGAACATCGTGAAGGGCAT
GAGAGAGCTCCGGGAGGTGCTGCGGACTGTGGAGACCAGCAACTCAGAACTTCAAAGTGATGGCGGCCAAGCA
CCTGGCGGGGTCTGCTGCACCTCCTGAGTGAGGAGTGCTACTGGAGCCCCCTGTCCACCCTCTGCCTGAGTT
CATGGGCAAGGAGGAGAGTTCCTTTCCGCACTCAGGCCCTGCGGAAACCTCACCTCTATGAAGGAGACAACCTCTA
CTGCCCCAAGGACAACATCGAGGAAGCCCTCCTGCTCCTCATCAGCGAATCCATGGCAACTCGAGATGTGGT
GCTGAGCCGGGTGCGGAGCAGGAGGAGGACCGGACAGTGAGCTTGCAGAATGCCGAGCCATCTATGACCTCCT
GAGCATCACGTTGGGCAGAAGGGGACAGTACGTCATGCTCTCGGAGTGCTGAGCGAGCCATGAAGTTTGCGTT
TGGAGAATTTACCTTTGGTACCAGGTGGCCCTCTCCATGGTGGCTTGTGGGAAGTCAGCCTACGCTGTGTCCCT
GCTGCGGGAGTGTTGAAGTTGCGGCCCTCGGACCCACCGTGCCCTGATGGCCGCGAAGGTCTGCATCGGGTC
CCTTCGCTGGCTAGAGGAAGCAGAGCACTTTGCCATGATGGTGATCAGCCTCGGAGAGGAAGCCGGGAGTTCTCT
CCCCAAGGGCTACCTGGCTCTGGGTCTCACCTATAGCCTGCAGGCCACCGACGCCACCCTGAAGTCCAAGCAAGA
TGAATTGCACCGGAAGGCACTGCAGACGCTGGAGAGGGCTCAGCAGCTGGCGCCAGTGACCCAGGTATCCT
CTATGTCTCGCTGCAGCTGGCCCTCGTCCGACAGATCTCCAGTGCCATGGAGCAGCTGCAGGAGCCCTGAAGGT
ACGCAAGGATGATGCCACGCCCCTCCACCTGCTGGCACTGCTCTTCTCTGCCAGAGCACCACGAGCATGCCCT
GGATGTTGTCAACATGGCCATCACCGAGCACCCCTGAGAACTTCAACCTGATGTTTACCAAGGTGAAGCTGGAGCA
GGTGCTGAAAGGCCAGAGGAAGCCCTCGTGACCTGCAGACAAGTGCTGAGGCTGTGGCAGACCCTGTACAGCTT
CTCCAGCTGGGAGGCCTAGAAAAGGATGGCAGCTTCGGTGAGGGCCTACCATGAAGAAGCAGAGTGGCATGCA
CCTGACTTTGCCTGATGCCATGATGCAGACTCTGGCTCCCGCGGGCTTCGTCCATCGCCGCTCCCGGCTGGA
GGAGGCCATGTGAGAGCTGACTATGCCCTCTTCGGTCTGAGCAGGGCCCCATGCAGCTGTGGACCAGCTGGA
ACAGATCTGGCTGCAGGCTGCTGAGCTGTTATGAGGAGCAGCAGCCTCAAGGAAGCAGGTTTCTGCATCCAGGA
GGCGGGGGGCTCTTCCCACTTCTCACTCAGTACTCTATATGCGGGGCGGGCTGGCTGAGGTGAAGGGCAACCT
GGAGGAGGCCAAGCAGCTGTACAAGGAGGCGCTCACGGTGAAACCAGATGGCGTGCGCATCATGCATAGCCTGGG
TCTGATGCTGAGTCGGCTGGGCCACAAGAGCTTGCGCCAGAAGGTGCTTCGTGATGCCGTGGAGAGGCAGAGTAC
GTGCCACGAGGCGTGCGAGGCTGGGCGAGGTGCTGCAGGCCAGGGCCAGAACGAGGCTGCCGTTGACTGCTT
CCTCACGCCCCCTTGAGCTGAGGCGCAGCAGCCTGTACTGCCCTTCTCCATCATCCCCAGAGAGCTCTGACGACG
CTGCAGCCGAGGAGGGGCTGGCCAGAGGGAGAGGCAGCAGGGAACGTGGGTGAGGTTGGGCAACAGTG
GCATCAGGTGCGGGGCTCAGGGAAATACATCTTTAGTGAACGCCCTCTGCAGCTGCAGCCCTCGTTCTCTGGCT
GGGCCAAGAGGGGCTTCCCTGGATTTCTTTGTTGGTGCTTGGGAAACAGTCTGACTTGAACCCTAAGTGCCCTTG
GAGAGTTTTGTGGTGACCAGACTTGCTCCCCAAGAGCTGGGCAGCGGGAGCCTCACAGCTGTCTTACCCTCA
CCCATGCCCTCTGGCTTGGAGTCTGGGTGGGGGTTCTCACTCCCCACTCTCAGCACAGTACAGACTTCTGGATCT
CTCTCAGGTCTTGCCAGGGCGGTACAATGTGAAGAACTGCGGGCAAGTGGAAGACTATGAGATTTCTGGGT
TCCCTTCTCAGACTTGGAGTTAGTAGATGATTCCTGCATTGCCCTGCTTGCCCTCTGAGACCAGCTGGGCCCCA
CCTTGCTCTTTCCCCCTGCTACCAAGTGCCCTTTGGGGTCTGACCAGGGGTACTGAGCACCGGCCCTAACACTTCC
ATCTCCACCCACCCCATCTCCCTGGCGATGTGCTCCAGCCCAAGCAGCCTCCGTAGGCTTTAGATCCTGTGGTTG
CTAGATCCAGTCCCTTTCTAATACCCTGAGTCAACACATTACTCTGCAGGTCTTAGGCTACAATGCAGGTCCCTT
GAGGGCCACCAACATGGAGGTAGGCAGTTTCTAGGACTGTCCCCAGTACATCTCACCACCCACAGCCCTTTTTT
GCCTTGATTGAGCCCTACCCCTGGCCTTTTGGCTTCCCTGCCTGAGAGAGACCTGAGGAGGGGACAGAGCCAG
CCCCCTCTCCTGTGGCTGAGCAGGCCTCTGTGTCCATGACACCTGTCTTCCGGGCTGGGGGCTGTGGGTGTATGT
CCTCCCTACTGGCTTCCCCGGCCCCCTGCTGCATGATGCTCTTGGAACCTCTTCTCCAAGGAGTCAGTCCCCAGGC
CTATCAGGGGATCCTTTTGTATCTGCACTTTGGGTTTTAGTTTTCAAAGCTCCATCAGGTACAGCTTGCAATTCAG
GATGTGTGAAAGCTCGGGTGAGGGCTGCCCTGGTTCATCATAGCTCCACCTTCTCGGAAGGAGTGGGCTGTG

791/6881
FIGURE 738B

GAGACCCCCCATCCATGGCACACTAGCTCAGCACTGCATTTCCCGAGATGATTCCCAAGACAGCTGGTGCCTCCT
GGCTTTTCTGTGCCAGGCCAAGGGGCACCACAGAGGACCCTGGATCCTTTGCCTCTTCTTGGTGAAGGATCTCT
ATGTATGTGTATATAAATATAGTTTTTTATCTATATATAAAATAGAGATCTATTTTTTTCTGGAATTCTG
TTAGAAAAGTAAAGAAAAAGCAAATGCTGTTGGTTTATCTCAGGGTGCCCAAAGTGGTTATAGTCAATTTTGGT
ACTAGGAAAGGCACCCAATGCATTTCTGACTTTTAAGCATTTCCTTGTTGGAAGCAGCAGAGGGCCAGGCCAAG
TTGCTGACAGTGACTTTGCAGGTTGAATAAAGAAACCTTGGAGGGGAAGCAGGCTTGTCTGAAGCAGCATGTAT
ATTCACTGGGCATGTAGCTCCACACCAGCCTTGAGCCAGGCCCTGGACAGGAGGGGCTGTTGCAGGATGAGGGA
GGCCAGAGAAGGCATCGAAGCCAAGACCTGGGCCCACCTGGGGAGGGATGTGGGAAAGGAAGGATGGGAGGGAGG
ACCTCTGGGAAAATGTGGATTGAGCTGGTGAGAGTGTGCTAAGGCTGGGCTAAAGCCTGGAGAGGGTAGGAG
GAGGCAAGAGGGGTCCAGGCAGGGCTGATCCTGGCCTCTGACCTGTCCAGGGCGACCCCTGAAGCCCCTGCTGCC
TCTGGGCATTGCTGGGAGAGGCCAAGGCAGGACTCAGTCTGAACAGAGATCCCCTCGGGCATTGCTGATGGGCC
ACCTTCAGCTGCAGGGAAGAAGCCTAGGAGAGGAGGCATGGGAGGGACCTGGGCCTTGTTTCAGATTGGCCACCTC
TGCTGAGAAGTCCATACCAAGTACCCCCCTAATAAGTTATGCCACATACCAACGTACTGTGGATATTATAACCTGC
ATTAAACAACCTCTAAAGAACGCTGCTCATT

792/6881
FIGURE 739

MAAKGAHGSYLKVESELERCRAEGHWDRMPELVRQLQTLSPG GGGNRRGSPSAAFTFPD TDDFGKLLLAEEALLE
QCLKENHAKIKDSMPLEKNEPKMSEAKNYLSSILNHGRLSPQYMCEAMLILGKLHYVEGSYRDAISMYARAGID
DMSMENKPLYQMRLLSEAFVIKGLSLERLPNSIASRFLTEREEEVITCFERASWIAQVFLQELEKTINNSTSRH
LKGCHPLDYELTYFLEAALQSAYVKNLKKGNIVKGMRELREVLRTVETKATQNFKVMAAKHLAGVLLHSLSEECY
WSPLSHPLPEFMGKEESSFATQALRKPHLYEGDNLYCPKDNIEEALLLLISESMATR DVVLSRVPEQEEDRTVS
LQNAAIYDLLSITLGRRGQYVMLSECLERAMKFAGFEHLWYQVALSMVACGKSAYAVSLLRECVKLRPSDPTV
PLMAAKVCIGSLRWLEEAHFAMMVISLGEEAGEFLPKGYLALGLTYSLQATDATLKSQDELHRKALQTLERAQ
QLAPSDPQVILYVSLQALVRQISSAMEQLQEALKVRKDDAHALHLLALLFSAQKHHQHALDVVNMAITEHPENF
NLMFTKV KLEQVLKGPEEALVTCRQVLR LWQTLYSFSQLGGLEKDGSFGEGLTMKKQSGMHLTL PDAHDADSGSR
RASSIAASRLEEAMSELTMPSSVLKQGPMQLWTTLEQIWLQAAELFMEQQHLKEAGFCIQEAAGLFTSHSVLYM
RGR LAEVKGNLEEAKQLYKEALT VNP DGVRIMHSLGLMLSRLG HKSLAQKVL RDAVERQSTCHEAWQGLGEVLQA
QGQNEAAVDCFLTALELEASSPVLPFSIIPREL

793/6881
FIGURE 740

AGTCCGAGTGGAGAGAGCGAGCTGAGTGGTTGTGTGGTTCGCGTCTCGGAAACCGGTAGCGCTTGCAGCATGGCTG
ACCAACTGACTGAAGAGCAGATTGCAGAATTCAAAGAAGCTTTTTCACTATTGACAAAGATGGTGATGGAACATA
TAACAACAAAGGAATTGGGAACTGTAATGAGATCTCTTGGGCAGAATCCACAGAAGCAGAGTTACAGGACATGA
TTAATGAAGTAGATGCTGATGGTAATGGCACAATTGACTTCCCTGAATTTCTGACAATGATGGCAAGAAAAATGA
AAGACACAGACAGTGAAGAAGAAATTAGAGAAGCATTCCGTGTGTTTGATAAGGATGGCAATGGCTATATTAGTG
CTGCAGAACTTCGCCATGTGATGACAAACCTTGGAGAGAAGTTAACAGATGAAGAAGTTGATGAAATGATCAGGG
AAGCAGATATTGATGGTGATGGTCAAGTAACTATGAAGAGTTTGTACAAATGATGACAGCAAAGTGAAGACCTT
GTACAGAATGTGTTAAATTTCTTGTACAAAATTGTTTATTTGCCTTTTCTTTGTTTGTAACCTTATCTGTAAAAGG
TTTCTCCCTACTGTCAAAAAAATATGCATGTATAGTAATTAGGACTTCATTCTCCATGTTTTCTTCCCTTATCT
TACTGTCAATTGTCCTAAAACCTTATTTTAGAAAATTGATCAAGTAACATGTTGCATGTGGCTTACTCTGGATATA
TCTAAGCCCTTCTGCACATCTAACTTAGATGGAGTTGGTCAAATGAGGGAACATCTGGGTTATGCCTTTTTTAA
AGTAGTTTTCTTTAGGAACTGTCAGCATGTTGTTGTTGAAGTGTGGAGTTGTAACCTCTGCGTGGACTATGGACAG
TCAACAATATGTACTTAAAAGTTGCACTATTGCAAAACGGGTGATTATCCAGGTACTCGTACACTATTTTTTTG
TACTGCTGGTCCTGTACCAGAAACATTTCTTTTATTGTTACTTGCTTTTTTAACTTTGTTTAGCCACTTAAAAAT
CTGCTTATGGCACAATTTGCCTCAAAATCCATTCCAAGTTGTATATTGTTTTCCAATAAAAAAATTACAATTT

794/6881
FIGURE 741

GGCGGGAAACAGCTTAGTGGGTGTGGGGTCGCGCATTTTCTCAACCAGGAGGTGAGGAGGTTTCGACATGGCGG
TGCAGCCGAAGGAGACGCTGCAGTTGGAGAGCGCGGCCGAGGTGGCTTCGTGCGCTTCTTTCAGGGCATGCCGG
AGAAGCCGACCACCACAGTGCGCCTTTTCGACCGGGGCGACTTCTATACGGCGCACGGCGAGGACGCGCTGCTGG
CCGCCCCGGGAGGTGTTCAAGACCCAGGGGGTGATCAAGTACATGGGGCCGGCAGGAGCAAAGAATCTGCAGAGTG
TTGTGCTTAGTAAATGAATTTTGAATCTTTGTAAAAGATCTTCTTCTGGTTCGTGAGTATAGAGTTGAAGTTT
ATAAGAATAGAGCTGGAAATAAGGCATCCAAGGAGAATGATTGGTATTGGCATATAAGGCTTCTCCTGGCAATC
TCTCTCAGTTTGAAGACATTCTCTTTGGTAACAATGATATGTCAGCTTCCATTGGTGTGTGGGTGTTAAATGT
CCGCAGTTGATGGCCAGAGACAGGTGGAGTTGGGTATGTGGATTCCATACAGAGGAACTAGGACTGTGTGAAT
TCCCTGATAATGATCAGTTCTCCAATCTTGAGGCTCTCCTCATCCAGATTGGACCAAAGGAATGTGTTTACCCG
GAGGAGAGACTGCTGGAGACATGGGGAACTGAGACAGATAATTCAAAGAGGAGGAATCTGATCACAGAAAGAA
AAAAAGCTGACTTTTCCACAAAAGACATTTATCAGGACCTCAACCGTTGTTGAAAGGCAAAAAGGGAGAGCAGA
TGAATAGTGCTGTATTGCCAGAAATGGAGAATCAGGTTGCAGTTTCATCACTGTCTGCGGTAATCAAGTTTITAG
AACTCTTATCAGATGATTCCAATTTGGACAGTTTGAAGTACTACTTTTGAAGTTCAGCCAGTATATGAAATTTGG
ATATTGCAGCAGTCAGAGCCCTTAACCTTTTTCAGGGTCTGTTGAAGATACCACTGGCTCTCAGTCTCTGGCTG
CCTTGCTGAATAAGTGTAACCCCTCAAGGACAAAGACTTGTAAACAGTGGATTAAAGCAGCCTCTCATGGATA
AGAACAGAATAGAGGAGAGATTGAATTTAGTGGAGCTTTTGTAGAAGATGCAGAATTGAGGCAGACTTTACAAG
AAGATTTACTTCGTGCGATTCCAGATCTTAACCGACTTGCCAAGAAGTTTCAAAGACAAGCAGCAAATTTACAAG
ATTGTTACCGACTCTATCAGGGTATAAATCAACTACCTAATGTTATACAGGCTCTGGAAAAACATGAAGGAAAAAC
ACCAGAAATTTGTTGGCAGTTTGTGACTCCTCTTACTGATCTTCGTCTGACTTCTCCAAGTTTTCAGGAAA
TGATAGAAACAACTTTAGATATGGATCAGGTGGAAAACCATGAATTCCTTGTAACCTTCATTTGATCCTAATC
TCAGTGAATTAAGAGAAATAATGAATGACTTGGAAAAGAAGATGCAGTCAACATTAATAAGTGCAGCCAGAGATC
TTGGCTTGGACCTTGGCAAACAGATTAACTGGATTCCAGTGCACAGTTTGGATATTACTTTTCGTGTAACCTGTA
AGGAAGAAAAAGTCTTCGTAACAATAAAACTTTAGTACTGTAGATATCCAGAAGAATGGTGTTAAATTTACCA
ACAGCAAATGACTTCTTTAAATGAAGAGTATACCAAAAATAAAACAGAAATATGAAGAAGCCAGGATGCCATTG
TTAAAGAAATGTCAATATTTCTTCAGGCTATGTAGAACCAATGCAGACACTCAATGATGTGTAGCTCAGCTAG
ATGCTGTTGTGCTGCTTGTGCTCAGGTGTCAAATGGAGCACCTGTTCCATATGTACGACCAGCCATTTTGGAGAAAG
GACAAGGAAGAATTATATTAAGCATCCAGGCATGCTTGTGTTGAAGTTCAAGATGAAATTCATTTATTCCTA
ATGACGTATACCTTGAAAAAGATAAACAGATGTTCCACATCATTACTGGCCCAATATGGGAGGTAAATCAACAT
ATATTGCAAACTGGGGTGATAGTACTCATGGCCCAATTTGGGTGTTTTGTGCCATGTGAGTCAGCAGAAGTGT
CCATTGTGGACTGCATCTTAGCCCGAGTAGGGGCTGGTGACAGTCAATTGAAAGGAGTCTCCACGTTTCATGGCTG
AAATGTTGGAACTGCTTCTATCCTCAGGTCTGCAACCAAAGATTCAATTAATAATCATAGATGAATTGGGAAGAG
GAACCTTCTACCTACGATGGATTGGGTAGCATGGGCTATATCAGAATACATTGCAACAAAGATTGGTGCTTTTT
GCATGTTTGCAACCAATTTTCATGAACCTTACTGCCTTGCCAATCAGATACCAACTGTTAATAATCTACATGTCA
CAGCACTCACCCTGAAGAGACCTTAACCTATGCTTTATCAGGTGAAGAAAGGTGTCTGTGATCAAAGTTTGGGA
TTCATGTTGCAGAGCTTGCTAATTTCCCTAAGCATGTAATAGAGTGTGCTAAACAGAAAGCCCTGGAACCTGAGG
AGTTTCAGTATATTGGAGAATCGCAAGGATATGATATCATGGAACAGCAGCAAAGAAGTGCTATCTGGAAGAG
AGCAAGGTGAAAAAATTATTTCAGGAGTTCTGTCCAAGGTGAAACAAATGCCCTTTACTGAAATGTCAGAAGAAA
ACATCACAATAAAGTTAAACAGCTAAAGCTGAAGTAATAGCAAAGAATAATAGCTTTGTAAATGAAATCATT
CACGAATAAAAGTTACTACGTGAAAAATCCAGTAATGGAATGAAGGTAATATTGATAAGCTATTGTCTGTAATA
GTTTTATATTGTTTTATATTAACCTTTTCCATAGTGTTAACTGTGAGTGGCCATGGGCTATCACTTAATAAG
ATATTAGTAATATTTTACCTTTGAGGACATTTTCAAAGATTTTATTTTGAAGAAATGAGAGCTGTAAGTGGAC
TGTTTGAATTGACATAGGCAATAATAAGTGATGTGCTGAATTTTATAAATAAAATCATGTAGTTTGTGG

795/6881
FIGURE 742

MAVQPKETLQLESAAEVGFVRFFQGMPEKPTTTVRLFDRGDFYTAHGEDALLAAREVFKTQGVIKYMGPAGAKNL
QSVVLSKMNFEFVKDLLLVQRQYRVEVYKNRAGNKASKENDWYLAYKASPGNLSQFEDIILFGNNDMSASIGVGV
KMSAVDQGRQVGVGYVDSIQKRLGLCEFPDNDQFSNLEALLIQIGPKCVLPGETAGDMGKLRQIIQRGGILIT
ERKKADFSTKDIYQDLNRLKGGKGEQMNSAVLPemenQVAVSSLSAVIKFLELLSDDSNFGQFELTTDFDSQYM
KLDIAAVRALNLFQGSVEDTTGSQSLAALLNKCKTPQGQRLVNQWIKQPLMDKNRIEERLNLVEAFVEDAELRQT
LQEDLLRRFPDLNRLAKKFQRQAANLQDCYRLYQGINQLPNVIALEKHEGKHQKLLAVFVTPLTDLRSDFSKE
QEMIETTLDMQVENHEFLVKPSFDPNLSELREIMNDLEKKMQSTLISAARDLGLDPGKQIKLDSSAQFGYYFRV
TCKEEKVLRNNKNFSTVDIQKNGVKFTNSKLTSLNEEYTKNKTEYEEAQDAIVKEIVNISSGYVEPMQTLNDVLA
QLDAVVSFAHVSNGAPVYVRPAILEKGQGRILKASRHACVEVQDEIAFIPNDVYFEKDKQMFHIITGPNMGGK
STYIRQTGVIVLMAQIGCFVPCESAIVDCILARVGAGDSQLKGVSTFMAEMLETASILRSATKDSLIIIDEL
GRGTSTYDGFGLAWAISEYIATKIGAFCMFATHFHELTALANQIPTVNNLHVTALTTEETLTMLYQVKKGVCDQS
FGIHVAELANFPKHVIECAKQKALELEEFQYIGESQGYDIMEPAAKKCYLEREQGEKIIQEFLSKVKQMPFTEMS
EENITIKLKQLKAEVIAKNNSFVNEIISRIKVT

796/6881

FIGURE 743

CTTTTGCGGGTGGCGGCGAACGCGGAGAGCACGCCATGAAGGCCTCGGGCACACTACGAGAGTACAAAGTAGTGG
GTCGCTGCCTGCCCACCCCAAATGCCACACACCACCCCTCTACCGCATGCGAATCTTTGCGCCTAATCATGTCG
TCGCCAAGTCCCGCTTCTGGTACTTCGTATCTCAGTTAAAGAAGATGAAGAAGTCTTCAGGGGAGATTGTCTACT
GTGGGCAGGTGTTTGAGAAGTTCCCCTTGCGGGTGAAGAACTTCGGGATCTGGCTGCGCTATGACTCCCGGAGCG
GCACCCACAACATGTACCGGAATACCGGGACCTGACCACCGCAGGCGCTGTCACCCGGTGCTACCGAGACATGA
GCGCCCGGCACGGCGGCCGGGCCCCACTCCATTCAGATCATGAAGGTGGAGGAGATCGCGGCCAGCAAGTGCCGCC
GGCCGGCTGTCAAGCAGTTCCACGACTCCAAGATCAAGTTCCCGCTGCCCCACCGGGTCTGCGCCGTCAGCACA
AGCCACGCTTCACCACCAAGAGGCCCAACACCTTCTTCTAGGTGCAGGGCCCTCGCCCGGTGTGCCCCAAATAA
ACTCAGGAACGCCCC

797/6881
FIGURE 744A

ATTTCGCCGCCAGCAGGAGCCGCGCGGTAGATGCGGTGCTTTTAGGAGCTCCGTCCGACAGAACGGTTGGGCCTTG
CCGGCTGTCCGTATGTCGCGACAGAGCACCCTGTACAGCTTCTTCCCCAAGTCTCCGGCGCTGAGTGATGCCAAC
AAGGCCTCGGCCAGGGCCTCACGCGAAGGCGGCCGTGCGGCCGTGCCCCGGGGCCTCTCCTTCCCCAGGCGGG
GATGCGGCCCTGGAGCGAGGCTGGGCCTGGGCCCAGGCCCTTGGCGCGATCCGCGTCACCGCCCAAGGCGAAGAAC
CTCAACGGAGGGCTGCGGAGATCGGTAGCGCCTGCTGCCCCACCAGTTGTGACTTCTCACCAGGAGATTGGTT
TGGGCCAAGATGGAGGGTTACCCCTGGTGGCCTTGTCTGGTTTACAACCACCCCTTTGATGGAACATTTCATCCGC
GAGAAAGGGAAATCAGTCCGTGTTTCATGTACAGTTTTTGTATGACAGCCCAACAAGGGGCTGGGTTAGCAAAAGG
CTTTTAAAGCCATATACAGGTTCAAATCAAAGGAAGCCAGAAGGGAGGTCATTTTACAGTGCAAAGCCTGAA
ATACTGAGAGCAATGCAACGTGCAGATGAAGCCTTAAATAAAGACAAGATTAAGAGGCTTGAATTGGCAGTTTGT
GATGAGCCCTCAGAGCCAGAAGAGGAAGAAGAGATGGAGGTAGGCACAACCTTACGTAACAGATAAGAGTGAAGAA
GATAATGAAATTGAGAGTGAAGAGGAAGTACAGCCTAAGACACAAGGATCTAGGCGAAGTAGCCGCCAAATAAAA
AAACGAAGGGTCATATCAGATTCTGAGAGTGACATTGGTGGCTCTGATGTGGAATTTAAGCCAGACACTAAGGAG
GAAGGAAGCAGTGATGAAATAAGCAGTGGAGTGGGGGATAGTGAGAGTGAAGGCCTGAACAGCCCTGTCAAAGTT
GCTCGAAAGCGGAAGAGAATGGTGACTGGAAATGGCTCTCTTAAAAGGAAAAGCTCTAGGAAGGAAACGCCCTCA
GCCACCAAAACAAGCAACTAGCATTTCATCAGAAACCAAGAATACTTTGAGAGCTTTCTCTGCCCTCAAATTTCT
GAATCCCAAGCCCACGTTAGTGGAGGTGGTGATGACAGTAGTCGCCCTACTGTTTGGTATCATGAAACTTTAGAA
TGGCTTAAGGAGGAAAAGAGAAGAGATGAGCACAGGAGGAGGCCTGATCACCCCGATTTTGATGCATCTACACTC
TATGTGCCTGAGGATTTCTCAATTCTTGTACTCCTGGGATGAGGAAGTGGTGGCAGATTAAGTCTCAGAACTT
GATCTTGTCTATCTGTTACAAGGTGGGGAATTTTATGAGCTGTACCACATGGATGCTCTTATTGGAGTCAGTGAA
CTGGGGCTGGTATTTCATGAAAGGCAACTGGGCCCATTTCTGGCTTTCTGAAATTGCAATTTGGCCGTTATTTCAGAT
TCCCTGGTGCAGAAGGGCTATAAAGTAGCACAGTGGAAACAGACTGAGACTCCAGAAATGATGGAGGCACGATGT
AGAAAGATGGCACATATATCCAAGTATGATAGAGTGGTGAAGGAGGAGATCTGTAGGATCATTACCAAGGGTACA
CAGACTTACAGTGTGCTGGAAGGTGATCCCTCTGAGAACTACAGTAAGTATCTTCTTAGCCCTCAAAGAAAAAGAG
GAAGATTCTTCTGGCCATACTCGTGATATGGTGTGTGCTTTGTTGATACTTCACTGGGAAAGTTTTTCATAGGT
CAGTTTTTCAGATGATCGCCATTGTTTCGAGATTTAGGACTCTAGTGGCACACTATCCCCAGTACAAGTTTTATTT
GAAAAAGGAAATCTCTCAAAGGAACTAAAACAATTCTAAAGAGTTTCATTGCTCTCTTTCAGGAAGGTCTG
ATACCCGGCTCCCAAGTTTTGGGATGCATCCAAAGCTTTGAGAACTCTCCTGAGGAAGAATATTTTAGGGAAAAG
CTAAGTGATGGCATTGGGGTGATGTTACCCAGGTGCTTAAAGGTATGACTTCAGAGTCTGATTCCATTGGGTTG
ACACCAGGAGAGAAAAGTGAATTGGCCCTCTCTGCTCTAGGTGGTTGTGCTTCTACCTCAAAAAATGCCTTATT
GATCAGGAGCTTTTATCAATGGCTAATTTGAAGAATATATTCCTTGGATTCTGACACAGTCAGCACTACAAGA
TCTGGTGCTATCTTACCAAAGCCTATCAACGAATGGTGTCTAGATGCAGTGACATTAAACAACCTTGAGATTTTT
CTGAATGGAACAAATGGTTCTACTGAAGGAACCCCTACTAGAGAGGGTTGATACTTGCCATACTCCTTTTGGTAAG
CGGCTCCTAAAGCAATGGCTTTGTGCCCCACTCTGTAACCATTATGCTATTAATGATCGTCTAGATGCCATAGAA
GACCTCATGGTTGTGCCTGACAAATCTCCGAAGTTGTAGAGCTTCTAAAGAAGCTTCCAGATCTTGAGAGGCTA
CTCAGTAAAATTCTAATGTTGGGTCTCCCTGAAGAGTCAGAACCACCCAGACAGCAGGGCTATAATGTATGAA
GAACTACATACAGCAAGAAGATTATTGATTTTTCTTTCTGCTCTGGAAGGATTCAAAGTAATGTGTAAAATT
ATAGGGATCATGGAAGAAGTTGCTGATGGTTTTAAGTCTAAAATCCTTAAGCAGGTCATCTCTCTGCAGACAAAA
AATCCTGAAGGTCGTTTTCTGATTTGACTGTAGAATTGAACCGATGGGATACAGCCTTTGACCATGAAAAGGCT
CGAAAGACTGGACTTATTACTCCCAAAGCAGGCTTTGACTCTGATTATGACCAAGCTCTTGCTGACATAAGAGAA
AATGAACAGAGCCTCCTGGAATACCTAGAGAAAACAGCGCAACAGAATTGGCTGTAGGACCATAGTCTATTGGGGG
ATTGGTAGGAACCGTTACCAGCTGGAAATTCCTGAGAATTTCAACACTCGCAATTTGCCAGAAGAATACGAGTTG
AAATCTACCAAGAAGGGCTGTAAACGATACTGGACCAAACTATTGAAAAGAAGTTGGCTAATCTCATAAATGCT
GAAGAACGGAGGGATGATCATTGAAGGACTGCATGCGCGACTGTTCTATAACTTTGATAAAAATTACAAGGAC
TGGCAGTCTGCTGTAGAGTGATCGCAGTGTGGATGTTTTACTGTGCTGGCTAACTATAGTCAGGGGGTGAT
GGTCTATGTGTGCGCCAGTAATTCTGTTGCCGAAGATACCCCCCTTCTTAGAGCTTAAAGGATCACGCCAT
CCTTGCAATTACGAAGACTTTTTTTGGAGATGATTTTATTCCTAATGACATTCTAATAGGCTGTGAGGAAGAGGAG
CAGGAAAATGGCAAAGCCTATTGTGTGCTTGTACTGGACCAATATGGGGGGCAAGTCTACGCTTATGAGACAG
GCTGGCTTATTAGCTGTAATGGCCAGATGGGTGTACGTCCCTGCTGAAGTGTGCAGGCTCACACCAATTGAT

798/6881

FIGURE 744B

AGAGTGTCTTACTAGACTTGGTGCCTCAGACAGAATAATGTCAGGTGAAAGTACATTTTTTGTGGAATTAAGTGAA
ACTGCCAGCATACTCATGCATGCAACAGCACATTCTCTGGTGCTTGTGGATGAATTAGGAAGAGGTACTGCAACA
TTTGATGGGACGGCAATAGCAAATGCAGTTGTTAAAGAACTTGCTGAGACTATAAAATGTCGTACATTATTTTCA
ACTCACTACCATTTCATTAGTAGAAGATTATTCTCAAATGTTGCTGTGCGCCTAGGACATATGGCATGCATGGTA
GAAATGAATGTGAAGACCCAGCCAGGAGACTATTACGTTCTCTATAAATTCATTAAAGGGAGCTTGTCTTAA
AGCTATGGCTTTAATGCAGCAAGGCTTGCTAATCTCCAGAGGAAGTTATTCAAAGGGACATAGAAAAGCAAGA
GAATTTGAGAAGATGAATCAGTCACTACGATTATTTGGGAAGTTTGCCTGGCTAGTGAAAGGTCAACTGTAGAT
GCTGAAGCTGTCCATAAATTGCTGACTTTGATTAAAGGAATTATAGACTGACTACATTGGAAGCTTTGAGTTGACT
TCTGACCAAAGGTGGTAAATTCAGACAACATTATGATCTAATAAACTTTATTTTTTAAAAATGA

799/6881

FIGURE 745

MSRQSTLYSFFPKSPALSDANKASARASREGGAAAAAGASPSPGGDAWSEAGGPRPLARSASPPKAKNLNGG
LRRSVAPAAFTSCDFSPGDLVWAKMEGYPWPCLVYNHPFDGTFIREKGKSVRVHVQFFDDSPTRGWVSKRLKLP
YTGSKSKEAQKGGHFYSAPKPEILRAMQRADEALNKDKIKRLELAVCDEPSEPEEEEEEMEVTYVTDKSEEDNEI
ESEEVQPKTQGSRRSSRQIKRRVISDSSEDIGGSDVEFKPDTKEEGSSDEISSGVGDSESEGLNSPVKVARKR
KRMVTGNGSLKRKSSRKETPSATKQATSISSETKNTLRAFSAPQNSESQAHVSGGGDDSSRPTVWYHETLEWLKE
EKRRDEHRRRDPDFDASTLYVPEDFLNSCTPGMRKWWQIKSQNFDLVICYKVGKFYELYHMDALIGVSELGLV
FMKGNWAHSGFPEIAFGRYSDSLVQKGYKVARVEQTETPEMMEARCRKMAHISKYDRVVRREICRIITKGTQYTS
VLEGDPSENYSKYLLSLKEKEEDSSGHTRAYGVCVDTSLGKFFIGQFSDDRHCSTRFTLVAHYPPVQVLFKGN
LSKETKTIKSSLSLQEGSLIPGSQFWDASKTLRTLLEEEYFREKLSDGIGVMLPQVLKGMTSESDSIGLTPGE
KSELALSALGGCVFYLLKCLIDQELLSMANFEEYIPLSDTVSTTRSGAIFTKAYQRMVLDVTLNLEIFLNGT
NGSTEGTLLERVDTCHTPFGRLLKQWLCAPLCNHYAINDRLDAIEDLMVVPDKISEVVELLKKLPDLERLLSKI
HNVGSPKSNHPDSRAIMYEETYSKKKIIDFLSALEGFKVMCKIIGIMEEVADGFKSKILKQVISLQTKNPEG
RFPDLTVELNRWDATFDHEKARKTGLITPKAGFDSYDQALADIRENEQSLLEYLEKQNRNIGCRTIVYWGIGRN
RYQLEIPENFTTRNLPEEYELKSTKKGCKRYWTKTIEKKLANLINAEERRDVSLKDCMRRLFYNFDKNYKDWQSA
VECIAVLVDVLLCLANYSRGGDGPMCRPVILLPEDTPPFLELKGSRHPCITKTFFGDDFIPNDILIGCEEEEQENG
KAYCVLVTGPNMGKSTLMRQAGLLAVMAQMGCVPAEVCRLTPIDRVFTRLGASDRIMSGESTFFVELSETASI
LMHATAHSLVLVDELGRGTATFDGTAIANAVVKELAETIKCRTLSTHYHSLVEDYSQNVAVRLGHMACMVENEC
EDPSQETITFLYKFIKGACPKSYGFNAARLANLPEEVIQKGRKAREFEKMNQSLRLFREVCCLASERSTVDAAEAV
HKLLTLIKEL

800/6881
FIGURE 746A

GTGAGCTGAAGCAGGGCAGGGCATCAACTACCCAGGAAGTGCAAGGGGTTTGGGGATTTTCCTTTCCTAGCCAA
GGGAAGGCATGACAGACTGTACCTGGAAAAACAGGACACTCTTGCCCAAATACTGCACCTTTTGCACAGTCTTAG
CAACTGGCAGACCAGGAGATTCTCTCCTGTGCCTGATTCAATTGGGTCCCACACCCATAGGGCCTTGCTTACTGCC
AGTGCAGCAGTCTGAGATTAACACCCCATCCCCGGGAGAACTCTAAGAAGGAGCTGATGTGGAGGAGCAGCTGAG
ACAGTTCAAGATGACGACCACAGTAGCCACAGACTATGACAACATTGAGATCCAGCAGCAGTACAGTGATGTCAA
CAACCGCTGGGATGTGACGACTGGGACAATGAGAACAGCTCTGCGCGGCTTTTTGAGCGGTCCCGCATCAAGGC
TCTGGCAGATGAGCGTGAAGCCGTGCAGAAGAAGACCTTCACCAAGTGGGTCAATTCACCTTGCCCGTGTGTC
CTGCCGGATCACAGACCTGTACACTGACCTTCGAGATGGACGGATGCTCATCAAGCTGCTGGAGGTCTCTCTGG
AGAGAGGCTGCCATAACCCACCAAGGGACGAATGCGCATCCACTGCTTAGAGAATGTGGACAAGGCCCTTCAGTT
CCTGAAGGAGCAGAGAGTCCATCTTGAGAACATGGGGTCCCATGACATCGTGGATGGAAACCCCGGCTGACCCCT
TGGCCTCATCTGGACCATCATCTGCGCTTCAGATCCAGGATATCAGTGTGGAAACTGAAGACAACAAAGAGAA
GAAATCTGCCAAGGATGCATTGCTGTTGTGGTGCCAGATGAAGACAGCTGGGTACCCCAATGTCAACATTACAA
TTTACCACCTAGCTGGAGGGACGGCATGGCCTTCAATGCACTGATACACAACACCGGCCTGACCTGATAGATT
TGACAAACTAAAGAAATCTAACGCACACTACAACCTGCAGAATGCATTTAATCTGGCAGAACAGCACCTCGGCCT
CACTAAACTGTTGGACCCCGAAGACATCAGCGTGGACCATCCTGATGAGAAGTCCATAATCACTTATGTGGTGAC
TTATTACCACTACTTCTCTAAGATGAAGCCCTTAGCTGTTGAAGGAAAACGAATGGAAAGGTGCTTGACAATGC
TATTGAAACAGAAAAATGATTGAAAAGTATGAATCACTTGCCCTTGACCTTCTGGAATGGATTGAACAAACCAT
CATCATTCTGAACAATCGCAAATTTGCCAATTCAGTGGTGGGGTTCAACAGCAGCTTCAGGCATTCAACACTTA
CCGCACTGTGGAGAAACCAACCAAAATTTACTGAGAAGGGGAACTTGGAAGTGTGCTCTTCACCATTCAGAGCAA
GATGAGGGCCAAACAACCAAGGTCTACATGCCCCGGGAGGGGAAGCTCATCTCTGACATCAACAAGGCCTGGGA
AAGACTGGAAAAAGCGGAACACGAAAGAGAAGTGGCTTTGCGGAATGAGCTCATAAGACAGGAGAACTGGAACA
GCTCGCCCGCAGATTGATCGCAAGGCAGCTATGAGGGAGACTTGGCTGAGCGAAAACAGCGTCTGGTGTCTCA
GGACAACCTTTGGGTTTGACCTTCTGCAGTTGAGGCCGCCACAAAAAGCACGAGGCCATTGAGACAGACATTGC
CGCATACGAGGAGCGTGTGCAGGCTGTGGTAGCCGTGGCCAGGGAGCTCGAGGCCGAGAATTACCACGACATCAA
GCGCATCACAGCGAGGAAGGACAATGTCTATCCGGCTCTGGGAATACCTACTGGAAGTGTCTAGGGCCCGGAGACA
GCGGCTCGAGATGAACCTGGGGCTGCAGAAGATATTCAGGAAATGCTCTACATTATGGACTGGATGATGAAAT
GAAGGTGCTAGTATTGTCTCAAGACTATGGCAAACACTTACTTGGTGTGGAAGACCTGTTACAGAAGCACACCT
GGTTGAAGCAGACATTGGCATCCAGGCAGAGCGGGTGAGAGGTGTCAATGCCTCCGCCCAGAAGTTGCGAACAGA
CGGGGAAGGTTACAAGCCCTGTGACCCCAAGTGATCCGAGACCGCGTGGCCACATGGAGTTCTGTTATCAAGA
GCTTTGCCAGCTGGCGGCTGAGCGCAGGGCCCGTCTGGAAGAGTCCCGCCGCTCTGGAAGTTCTTCTGGGAGAT
GGCAGAAGAGGAAGGCTGGATACGGGAGAAGGAGAAGATCCTGTCTCGGACGATTACGGGAAAGACCTGACCAG
CGTCATGCGCCTGCTCAGCAAGCACCGGGCGTTCGAGGACGAGATGAGCGGCCGAGTGGCCACTTTGAGCAGGC
CATCAAGGAAGGCGAAGACATGATCGCGGAGGAGCACTTCGGGTGCGAGAAGATCCGTGAGAGGATCAATTACAT
CCGGGAGCAGTGGGCCAACCTAGAGCAGCTCTCGGCCATTGGAAGAAGCGCCTGGAGGAGGCCTCCCTGCTGCA
CCAGTTCAGGCAGATGCTGATGACATTGATGCCTGGATGCTGGACATCCTCAAGATTGTCTCCAGCAGCGACGT
GGGCCACGATGAGTATTCCACACAGTCTCTGGTCAAGAAACACAAGGACGTGGCGGAAGAGATCGCCAATTACAG
GCCCACCTTGACACGCTGCACGAACAAGCCAGCGCCCTCCCCAGGAGCATGCCGAGTCTCCAGACGTGAGGGG
CAGGCTGTGCGGCATCGAGGAGCGGTATAAGGAGGTGGCAGAGCTGACGCGGCTGCGGAAGCAGGCACTCCAGGA
CACTCTGGCCCTGTACAAGATGTTACGCGAGGCTGATGCCCTGTGAGCTCTGGATCGACGAGAAGGAGCAGTGGCT
CAACAACATGCAGATCCCAGAGAAGCTGGAGGATCTGGAGGTCAATCCAGCACAGATTGAGAGCCTAGAACCAGA
AATGAACAACAGGCTTCCCGGGTTGCAGTGGTGAACCAGATTGCACGCCAGCTGATGCACAGCGGCCACCCAAG
TGAGAAGGAAATCAAAGCCCAGCAGGACAACTCAACACAAGGTGGAGCCAGTTTCAAGAACTGAGGTTGACAGGAA
GAAGGATGCCCTCCTGTCTGCCCTGAGCATCCAGAACTACCCTCGAGTGCAATGAAACCAATCTGGATTGCG
GGAAAAGACCAAGGTCATCGAGTCCACCCAGGACCTGGGCAATGACCTGGCTGGCGTCATGGCCCTGCAGCGCAA
GCTGACCGGCATGGAGCGGACTTGGTGGCCATTGAGGCAAGCTGAGTGACCTGCAGAAGGAGGCGGAGAAGCT
GGAGTCCGAGCACCCCGACAGGCCAGGCCATCCTGTCTCGGCTGGCCGAGATCAGCGACGTGTGGGAGGAGAT
GAAGACCACCTTGAAAAACCGAGAGGCCTCCCTGGGAGAGGCCAGCAAGCTGCAGCAGTTCCTACGGGACTTGGA
CGACTTCAGTCTTGGCTCTCTAGGACCCAGACAGCGATCGCCTCGGAGGACATGCCAAACACCTGACCGAGGC

801/6881
FIGURE 746B

TGAGAAGCTGCTCACGCAGCACGAGAACATCAAGAATGAGATCGACAACACTACGAGGAGGACTACCAGAAGATGAG
GGACATGGGCGAGATGGTCACCCAGGGGCGAGACCGATGCCAGTACATGTTTCTGCGGCAGCGGCTGCAGGCCCT
GGACACTGGATGGAACGAGCTCCACAAGATGTGGGAGAACAGACAAAATCTCCTATCCCAGTCACATGCCATACCA
GCAGTTCCTCAGAGACACGAAGCAAGCCGAAGCCTTTCTTAACAACCAGGAGTATGTTCTGGCTCACACTGAAAT
GCCTACCACCTTGAAGGAGCTGAAGCAGCAATTAAAAAGCAAGAGGACTTCATGACCACCATGGACGCCAATGA
GGAGAAGATCAATGCTGTGGTGGAGACTGGCCGGAGGCTGGTGAGCGATGGGAACATCAACTCAGATCGCATCCA
GGAGAAGGTGGACTCTATTGATGACAGACATAGGAAGAATCGTGAGACAGCCAGTGAACTTTGATGAGGTTGAA
GGACAACAGGGATCTACAGAAATTCCTGCAAGATTGTCAAGAGCTGTCTCTCTGGATCAATGAGAAGATGCTCAC
AGCCCAGGACATGTCTTACGATGAAGCCAGAAATCTGCACAGTAAATGGTTGAAGCATCAAGCATTATGTCGAGA
ACTTGATCCAAACAAAGAATGGCTTGACAAAATCGAGAAGGAAGGAATGCAGCTCATTTCAGAAAAGCCTGAGAC
GGAAGCTGTGGTGAAGGAGAACTCACTGGTTTACATAAAATGTGGGAAGTCCTTGAATCCACTACCCAGACAAA
GGCCCAGCGGCTCTTTGATGCAAAACAGGCCGAACTTTTCACCCAGAGCTGTGCAGATCTAGACAAATGGCTGCA
CGGCTGGAGAGTCAGATTGATCTGATGACTATGGCAAACACCTGACCAGTGTCAATATCCTGCTGAAAAAGCA
ACAGATGCTGGAGAATCAGATGGAAGTGCGGAAGAAGGAGATCGAAGAGCTCCAAAGCCAGGCCAGGCCCTGAG
TCAGGAAGGGAAGAGCACCGACGAGGTAGACAGCAAGCGCTCACCCTGCAGACCAAGTTCATGGAGTTGCTGGA
GCCCTTGAACGAGAGGAAGCATAACCTGCTGGCCTCCAAAGAGATCCATCAGTTCAACAGGGATGTGGAGGACGA
GATCTGTGGGTTGGAGAGAGGATGCCCTTGGCAACTTCCACGGATCATGGCCACAACCTCCAGACTGTGCAGCT
GTTAATAAAGAAAAATCAGACCCTCCAGAAAGAAATCCAGGGGCACAGCCTCGCATTGACGACATCTTTGAGAG
GAGCCAAAACATCGTCACTGACAGCAGCAGCCTCAGCGCTGAGGCCATCAGACAGAGGCTTGCCGACCTGAAGCA
GCTGTGGGGTCTCCTCATTGAGGAGACAGAGAAACGCCACAGGCGGCTGGAGGAGGCGCACAGGGCCCAGCAGTA
CTACTTTGACGCTGCTGAGGCCGAAGCCTGGATGAGCGAGCAGGAGCTGTACATGATGTGAGAGGAGAAGGCCAA
GGATGAGCAGAGTGCTGTCTCCATGTTGAAGAAGCACCAGATCTTAGAACAGCTGTGGAGGACTATGCAGAGAC
CGTGATCAGCTCTCCAAGACCAGCCGGGCCCTGGTGGCCGACAGCCATCCTGAAAGTGAGCGCATTAGCAIGCG
GCAGTCCAAAGTGGATAAACTGTACGCTGGTCTGAAAGACCTTGCTGAAGAGAGAAGAGGCAAGCTGGATGAGAG
ACACAGGTTATTCCAGCTCAACCGGGAGGTGGACGACCTGGAGCAGTGGATCGCTGAGAGGGAGGTGGTTCGAGG
GTCCCATGAAC TGGGACAGGACTATGAGCATGTCACGATGTTACAAGAACGATTCCGGGAGTTTGCCCGAGACAC
CGGGAACATTGGGACAGGAGCGCGTGGACACGGTCAATCACCTGGCAGATGAGCTCATCACTCTGGACATTGAGA
TGCCGCCACCATCGCTGAATGGAAGGATGGCCTCAATGAAGCCTGGGCCGACCTCCTGGAGCTCATTGACACAAG
AACACAGATTCTTGCCGCTTCCTATGAACTGCACAAGTTTTACCACGATGCCAAGGAGATCTTTGGCGGTATACA
GGACAAACACAAGAACTCCCTGAGGAGCTTGGGAGAGATCAGAACACAGTGGAGACCTTACAGAGAATGCACAC
TACATTTGAGCATGACATCCAGGCTCTGGGCACACAGGTGAGGAGCTGCAGGAGATGCAGCCCGCTCCAGGC
GGCCTATGCGGGTGACAAGGCCGACGATATCCAGAAGCGCGAGAACGAGGTCCCTGGAAGCCTGGAAGTCCCTCCT
GGACGCTGTGAGAGCCGACGGGTGCGGCTGGTGGACACAGGGGACAAGTTCCGCTTCTTCAGCATGGTGCGGA
CCTCATGCTCTGGATGGAGGATGTCATCCGGCAGATCGAGGCCAGGAGAAGCCAAGGGATGTATCATCTGTTGA
ACTCTTAATGAATAATCATCAAGGCATCAAAGCTGAAATTGATGCACGTAATGACAGTTTACAACCTGCATTGA
ACTTGGGAAATCCCTGTTGGCGAGAAAACACTATGCATCTGAGGAGATCAAGGAAAAATTACTGCAGTTGACGGA
AAAGAGGAAAGAAATGATCGACAAGTGGGAAGACCGATGGGAATGGTTAAGACTGATTCTGGAGGTCCATCAGTT
CTCAAGAGACGCCAGTGTGGCCGAGGCTGGCTGCTTGGACAGGAGCCGTACCTATCCAGCCGAGAGATAGGCCA
GAGCGTGGACGAGGTGGAGAAGCTCATCAAGCGCCACGAGGCATTGAAAAGTCTGCAGCAACCTGGGATGAGAG
GTTCTCTGCCCTGGAAAGGCTGACTACATTGGAGTTACTGGAAGTGCGCAGACAGCAAGAGGAAGAGGAGAGGAA
GAGGCGGCCGCTTCTCCGAGCCGAGCACGAAGGTTTTCAGAGGAAGCCGAGTCCAGCAGCAGTGGGATACTTC
AAAAGGAGAACAAGTTTCCCAAACGGTTTGCCAGCTGAACAGGGATCTCCACGGATGGCAGAAACGGTGGACAC
AAGCGAAATGGTCAAGCGCGCTACAGAACAAGGACGAGCTCTAAAGAGTCCAGCCCCATCCCTCCCCGACCTC
TGATCGTAAAGCCAAGACTGCCCTCCAGCCAGAGTGCCGCCACCTTACCAGCCAGAACCAGGAGACACCTTC
GGCCAGATGGAAGGCTTCTCAATCGGAAACAGAGTGGGAGGCCCAATAAGAAAGCCTCAAGCAGGTCTCTG
GCACAATGTTTATTGTGTCTATAAATAACCAAGAAATGGGTTTCTACAAAGATGCAAGACTGCTGCTTCTGGAAT
TCCCTACCACAGCGAGGTCCCTGTGAGTTTGAAGAAGCTGTCTGCGAAGTGGCCCTTGATTACAAAAGAGAA
ACACGTATTCAAGCTAAGACTAAATGATGGCAATGAGTACCTCTTCCAAGCCAAAGACGATGAGGAAATGAACAC

802/6881

FIGURE 746C

ATGGATCCAGGCTATCTCTTCGCCATCTCCTCTGATAAACACGAGGTGTCTGCCAGCACCCAGAGCACGCCAGC
ATCCAGCCGCGCGCAGACCCCTCCCCACCAGCGTCGTCACCATCACCAGCGAGTCCAGTCCCGGCAAGCGGGAAAA
GGACAAAGAGAAAGACAAAGAGAAGCGGTTTCAGCCTTTTTGGCAAAAAGAAATGAATCCTTTCCTTCACCTCCT
GCCCTTCTCTTACCTTTTCAGTGAAATTCAGCATGCAAGCTCAGAACCAACACATTACTCTCTGTGCCTAATGT
TCCTCAATGTGGTTGATTATTTTTTTTTTTAATTTATAGAGCATTTCTGGGGGGGGTGGGG

803/6881
FIGURE 747

MTTIVATDYDNIEIQQQYSDVNNRWDVDDWDNENSSARLFERSRIKALADEREAVQKKTFTKWVNSHLARVSCRI
TDLYTDLRDGRMLIKLLEVLSEGERLPKPTKGRMRIHCLENVDKALQFLKEQRVHLENMGSHDIVDGNHRLTLGLI
WTIILRFQIQDISVETEDNKEKKSADALLLWCQMKTAGYPNVNIHNFITTSWRDGMFALIHKHRPDLIDFDKL
KKSNAHYNLQNAFNLAEQHLGLTKLLDPEDISVDHPDEKSIITYVVYYHYFSKMALAVEGKRIGKVLDNAIET
EKMIEKYESLASDLELWIEQTIIILNNRKFANSLVGVQQQLQAFNTYRTVEKPPKFTTEKGNLEVLLFTTIQSKMRA
NNQKVYMPREGKLISDINKAWERLEKAEHERELALRNELIRQEKLEQLARRFDRKAAMRETWLSNQRLVSQDNF
GFDLPAVEAATKKHEAIEDIAAYEERVQAVVAVARELEAENYHDIKRITARKDNVIRLWEYLLELLRARRQRLE
MNLGLQKIFQEMLYIMDWMDEMKVVLVSQDYGKHLGVEDLLQKHTLVEADIGIQAERVGVNASAQKFATDGEG
YKPCDPQVIRDRVAHMEFCYQELCQLAAERRARLEESRRLWKFFWEMAEWEGWIREKEKILSSDDYGKDLTSVMR
LLSKHRAFEDEMSEGRSGHFEQAIKEGEDMIAEEHFGESEKIRERIYYIREQWANLEQLSAIRKKRLEEASLLHQFQ
ADADDIDAWMLDILKIVSSSDVGHDEYSTQSLVKKHKDVAEEIANYRPTLDTLHEQASALPQEHAESEPDVRGRLS
GIEERYKEVAELTRLRKQALQDTLALYKMFSEADACELWIDEKEQWLNMMQIPEKLEDLEVIQHRFESLEPEMNN
QASRVAVVNIQARQLMHSGHPSEKEIKAQQDKLNTRWSQFRELVDKRDALLSALSQNYHLECNETKSWIREKT
KVIESTQDLGNDLAGVMALQRLTGEMERDLVAIEAKLSDLQKEAEKLESEHPDQAQAILSRALIESDVWEEMKTT
LKNREASLGEASKLQQFLRDLDDFQSWLSRTQTATASEDMPNTLTAEKLLTQHENIKNEIDNYEEDYQKMRDMG
EMVTQGGTDAQYMFRLRQLALDGTWNEHLHKMWENRQNLSSQSHAYQQFLRDTKQAEAFNNQYVLAHTEMPIT
LEGAEAAIKKQEDFMTTMDANEKINAVVETGRRLVSDGNINSRRIQEKVDSIDDRHRKNRETASELLMRLKDNR
DLQKFLQDCQELSLWINEKMLTAQDMSYDEARNLHSHKWLKHQAFMAELASNKEWLDKIEKEGMQLISEKPETEAV
VKEKLTGLHKMWEVLESTTQTKAQLRFDANKAELFTQSCADLDKWLHGLESQIQSDDYGKHLTSVNILLKKQQML
ENQMEVRKKEIEELQSQAQALSQEGKSTDEVDKRLTVQTKFMELLEPLNERKHNLASKEIHQFNRDVEDEILW
VGERMPLATSTDHGHNLQTVQLLIKKNQTLQKEIQGHQPRIDDIFERSQNIIVTDSSSLSAEAIQRLADLKQLWG
LLIEETEKRRHRLLEEAAHQYYFDAAEAEAWMSEQELYMMSEEKAKDEQSAVSMKKHQILEQAVEDYAEIVHQ
LSKTSRALVADSHPESESRISMRQSKVDKLYAGLKDIAEERRGKLDERHRLFQLNREVDDLEQWIAEREVVAGSHE
LGQDYEHVTMLQERFREFARDTGNIGQERVDVTNHLADELINSGHSDAATIAEWKDGLEAWADLLELIDTRTQI
LAASYELHKFYHDAKEIFGRIQDKHKKLPEELGRDQNTVETLQRMHTTFEHDIQALGTQVRQLQEDARLQAAAYA
GDKADDIQKRENEVLEAWKSLLDACESRRVRLVDTGDKFRFFSMVRDLMLWMDVIRQIEAQEKPRDVSSVELLM
NNHQGIKAEIDARNDSTTCIELGKSLARKHYASEEIKEKLLQLTEKRKEMIDKWEDRWELRLILEVHQFSRD
ASVAEAWLLGQEPYLSREIGQSVDEVEKLIKREAFKSAATWDERFSALERLTLELLEVRROQEEEEERKRRP
PSPEPSTKVSEEAESQQQWDTSKGEQVSQNGLPAEQGSFMAETVDTSEMVNGATEQRTSSKESPSPSTSDRK
AKTALPAQSAATLPAQTQETPSAQMEGFLNRKHEWEAHNKKASSRSWHNVYCVINNQEMGFYKDAKTAASGIPYH
SEVPVSLKEAVCEVALDYKKKKHVFKLRLNDGNEYLFQAKDDEEMNTWIAISSAISSDKHEVSASTQSTPASSR
AQLTPTSVVTITSESSPGKREKDKKDKKRFSLFGKKK

804/6881
FIGURE 748

CACAACCGCCCGCGGCTCTGAGACGCGGCCCCGGCGGCGGCGGCAGCAGCTGCAGCATCATCTCCACCCTCCAGC
CATGGAAGACGAGGAGGAAGAAGAGGAGGAGGAAGAGGAGGACGAGGACGAAGACCTGGAGGAGCTGGAGGTGCT
GGAGAGGAAGCCCGCCGCGGGCTGTCCGCGGCCCCAGTGCCCAACGCCCCCTGCCGCCGGCGCGCCCCTGATGGA
CTTCGGAATGACTTCGTGCCGCGCGCCCCGGGGACCCCTGCCGGCCGCTCCCCCGTCGCCCCGGAGCGGCA
GCCGTCTTGGGACCCGAGCCCGGTGTCGTGACCGTGCCCGCGCCATCCCCGCTGTCTGCTGCCGCAGTCTCGCC
CTCCAAGCTCCCTGAGGACGACGAGCCTCCGGCCCCG

805/6881
FIGURE 749

TTCTTCCTTTTCGATCCGCCATCTGCGGTGGAGCCGCCACCAAAATGCAGATTTTCGTGAAAACCCCTTACGGGGA
AGACCATCACCCCTCGAGGTTGAACCCTCGGATACGATAGAAAATGTAAAGGCCAAGATCCAGGATAAGGAAGGAA
TTCTTCCTGATCAGCAGAGACTGATCTTTGCTGGCAAGCAGCTGGAAGATGGACGTACTTTGTCTGACTACAATA
TTCAAAAGGAGTCTACTCTTCACTTTGTGTTGAGACTTCGTGGTGGTGCTAAGAAAAGGAAGAAGAAGTCTTACA
CCACTCCCAAGAAGAATAAGCACAAGAGAAAGAAGGTTAAGCTGGCTGTCCTGAAATATTATAAGGTGGATGAGA
ATGGCAAAATTAGTCGCCTTCGTCGAGAGTGCCCTTCTGATGAATGTGGTGCTGGGGTGTTTATGGCAAGTCACT
TTGACAGACATTATTGTGGCAAATGTTGTCTGACTTACTGTTTCAACAAACCAGAAGACAAGTAACTGTATGAGT
TAATAAAAGACATGAACTAACAAAAA

806/6881
FIGURE 750

MQIFVKLTGKTITLEVEPSDTIENVKAKIQDKEGIPPDQORLIFAGKQLEDGRTLSDYNIQKESTLHLVLRRLRG
GAKRKKKSYTTPKKNKHKKVLAFLKYYKVDENGKISRLRRECPSECGAGVFMASHFDRHYCGKCCLTYCF
NKPEDK

807/6881
FIGURE 751

GGCACGAGCAGAATCCAGGGGCCCGGGGCTGTAGATTCCTTGACAAGGATATCCTAGCGGCGAAACAACACCGTA
CTGGGAGTCAGAACGCTCTGGGTCTTAGTCTTGACTGCCATTAAGTAGCGGTATGACATTGGAGAAGCTTTTTTGA
CCCTTCTGGATTTCCGTTTCTTTCTGTAAAATGAGGAGCTTGGAAGATCCGGAAAATGAGGCCCATAGGAAAC
AAGTGACTTGCTGAGTCCAGATAAAGTACTGCTGACAGAGAGAAACATGAACCAGAAGCTACTGAAGTTGGAGAAC
TTGCTACGATTTACACTATTTATAGGCAACTGCACAGTCTGTGTCAAAGAAGAGCATTAAAGACAGTGGAGGCAT
GGGTTTTTCATCTGCTTACCCTGTGTGGACAGCTCAACTGTGTGCTGGCCCTGGCCAAACAGATGTGCTCAATGGG
GCTGCTTTATCTCAGTATAGGCTTCTAGTAACAAAAAGGAAGAAGGACCATGGAAATCTCAGTTATCTTCAACA
AAATCTAAAAAGGTGGTAGAAGTATGGATTGGAATGACTATTGAGGAACTGGCCAGGGCAATGGAAAAAACACA
GATTATGTATATGAAGCTTTATTGAACACTGATATTGACATAGATTCACTGGAAGCAGACTCACATTTAGATGAA
GTCTGGATCAAGAAGTGATAACGAAGGCAGGGATGAAGTTAAAGTGGAGTAAATTAAGACAGGACAAAGTCAGA
AAAAATAAAGATGCTGTAAGAAGGCCCCAGGCAGATCCAGCTTTATTAACCCCAAGGTCCCCAGTTGTTACTATA
ATGGGCCATGTTGATCACGGGAAAACGACATTACTTGACAAATTTGAAAACTCAAGTGGCAGCAGTGGAACT
GGAGGCATCACTCAGCACATTTGGTGCCTTTCTTGCTCTCTGCCTTCTGGGGAAAAGATAACTTTTCTTGATACT
CCAGGACATGCTGCTTTCTCAGCAATGAGAGCCAGAGGTGCTCAGGTCACTGACATTGTCGTATTGGTTGTAGCT
GCAGATGATGGAGTGATGAAACAACTGTAGAATCTATTGAGCATGCCAAAGATGCACAGGTTTCTATTATCCTT
GCCGTAATAAATGTGACAAAGCTGAGGCTGATCCTGAGAAAGTGAAAAAGAGCTGCTGGCTTACGATGTGGTA
TGTGAAGATTATGGAGGTGATGTTCAAGCAGTGCCCTGTCTCCGCACTTACGGGCGATAATCTGATGGCTTTGGCA
GAAGCAACAGTTGCTCTTGACAGAAATGTTAGAATTGAAAGCAGATCCCAATGGTCCAGTGGAAAGGAACAGTAATA
GAGTCTTTTACAGACAAAGGAAGAGGTCTTGTTACTACAGCTATAATTCAAAGAGGAACCTTAAGAAAAGGCTCT
GTTCTGGTTGCTGGAAATGTTGGGCAAAAGTACGCTTAATGTTTGATGAAATGGAAAAACAATTGATGAGGCC
TATCCCAGCATGCCAGTGGGAATTACAGGCTGGAGAGACCTTCTTCTGCAGGAGAAGAAATCTTGAAGTAGAA
TCTGAGCCAAGGGCACGTGAAGTTGTTGACTGGAGGAAATATGAACAAGAACAGGAGAAAGGTGAGGAGGATCTG
AAAATAATAGAAGAAAAGCGAAAGGAACACAAAGAAGCACATCAGAAAGCCCGTGAGAAGTATGGCCATCTACTG
TGGAAGAAGAGATCAATTCTACGGTTTTTAGAAAAGAAAAGAACAAATACCTTAAAGCCAAAAGAGAAAAGGGAA
AGAGATTCAAATGTACTTTCTGTGATTATTAAGGTGATGTTGATGGTTCTGTTGAGGCCATTTTGAACATTATA
GATACCTATGATGCTTCACACGAGTGTGAAGTGAATAGTACATTTTGGAGTGGGTGATATAAGTGCAAATGAT
GTTAACCTTGCTGAAACATTTGATGGTGTATATATGGCTTTAATGTGAATGCAGGCAATGTTATCCAACAGTCA
GCTGCAAAAAAGGAGTAAAAATTAACTTCACAAAATAATTTACCGTCTGTTGAAGATTTGCAAGAGGAACTG
AGCAGCAGATTACCCTGTGCTGTGGAAGAGCACCCAGTAGGTGAGGCATCTATACTAGCTACCTTCTCTGTAACA
GAAGGGAAGAAAAAGTTCTGTGGCTGGCTGCAGAGTCCAAAAGGGACAGTTAGAAAAACAAAAAAATTTAAA
CTAACCCGTAATGGACATGTAATTTGGAAGGGCTCATTACCTCATTGAAACACCATAAAGATGACATTTCAATT
GTCAAAACGGGAATGGATTGTGGTCTCAGTTTAGATGAAGACAATATGGAATTTCAAGTGGGAGACAGAATTGTT
TGTTATGAAGAAAAGCAAATTCAAGCCAAGACTTCTTGGGATCCAGGATTTTAAATTTACATTAAAAATGTAAAT
AACTCA

808/6881
FIGURE 752

MNQKLLKLENLLRFHTIYRQLHSLCQRRALRQWRHGFSSAYPVWTAQLCAWPWPTDVLNGAALSQYRLLVTKKEE
GPWKSQLSSTKSKKVVEVWIGMTIEELARAMEKNTDYVYEALLNTDIDIDSLEADSHLDEVWIKVITKAGMKLK
WSKLGQDKVRKNKDAVRRPQADPALLTPRSPVVTIMGHVDHGKTTLLDKFRKTQVAAVETGGITQHIGAFVLSLP
SGEKITFLDTPGHAAFSAMRARGAQVTDIVVLVVAADDGVMKQTVESIQHAKDAQVFIILAVNKCDKAEADPEKV
KKELLAYDVVCEDYGGDVQAVPVVSALTGDNLMALAEATVALAEMLELKADPNPVEGTVIESFTDKGRGLVTTAI
IQRGTLRKGSVLVAGKCWAKVRLMFDENGKTIDEAYPSMPVGITGWRDLPSAGEEILEVESEPRAREVVDWRKYE
QEQEKQEDLKIIEEKREKHEAHQKAREKYGHLLWKKRSILRFLERKEQIPLKPKEKRERDSNVLSVVIKGDVD
GSVEAILNIIDTYDASHECELELVHFGVDISANDVNLAETFDGVIYGFNVNAGNVIQQSAAKKGVKIKLHKIY
RLVEDLQEEELSSRLPCAVEEHPVGEASILATFSVTEGKKKVPVAGCRVQKGQLEKQKKFKLTRNGHVIWKGSLS
LKHKKDDISIVKTGMDCGLSLDEDNMEFQVGDRIVCYEEKQIQAKTSWDPGF

809/6881
FIGURE 753

CGGTGCCGGGTGTTCATGGCGGCTGCAGGTACTGCTGCTCGTGCCTCCGGCTCCGGCCCCCTGAGCGATGGTCCCTT
TCCTTCTGCCACGGCGGGATCGGGCACTCACCCAGTTGCAAGTGCGAGCACTATGGAGTAGCGCAGGGTCTCGAG
CTGTGGCCGTGGACTTAGGCAACAGGAAATTAGAAATATCTTCTGGAAAGCTGGCCAGATTTCAGATGGCTCTG
CTGTAGTACAGTCAGGTGACACTGCAGTAATGGTCACAGCGGTGAGTAAACAAACCTTCCCCTTCCAGTTTA
TGCCTTTGGTGGTTGACTACAGACAAAAAGCTGCTGCAGCAGGTAGAAATCCACAAACTATCTGAGAAGAGAGA
TTGGTACTTCTGATAAAGAAATTCTAACAAGTCGAATAATAGATCGTTCAATTAGACCGCTCTTTCAGCTGGCT
ACTTCTATGATACACAGGTTCTGTGTAATCTGTTAGCAGTAGATGGTGTAAATGAGCCTGATGCTTAGCAATTA
ATGGCGCTTCCGTAGCCCTCTCATTATCAGATATTCTTGGAAATGGACCTGTGGGGCAGTACGAATAGGAATAA
TTGATGGAGAATATGTTGTTAACCACAAGAAAAGAAATGTCTTCTAGTACTTTAAATTTAGTGGTTGCTGGAG
CACCTAAAAGTCAGATTGTTCATGTTGGAAGCCTCTGCAGAGAACATTTTACAGCAGGACTTTTGCCATGCTATCA
AAGTGGGAGTGAAATATACCCAACAAATAATTCAGGGCATTTCAGCAGTTGGTAAAAGAACTGGTGTACCAAGA
GGACACCTCAGAAGTATTTTACCCCTTCGCCAGAGATTGTGAAATATACTCATAAACTTGCTATGGAGAGACTCT
ATGCAGTTTTTACAGATTACGAGCATGACAAAGTTTTCCAGAGATGAAGCTGTTAACAAAATAAGATTAGATACGG
AGGAACAATAAAAAGAAAAATTTCCAGAAGCCGATCCATATGAAATAATAGAATCCTTCAATGTTGTTGCAAAGG
AAGTTTTTAGAAGTATTGTTTTGAATGAATACAAAAGGTGCGATGGTCGGGATTTGACTTCACCTAGGAATGTAA
GTTGTGAGGTAGATATGTTTAAACCCTTCATGGATCAGCATTATTTCAAAGAGGACAAACACAGGTGCTTTGTA
CCGTTACATTTGATTTCATTAGAATCTGGTATTAAGTCAGATCAAGTTATAACAGCTATAAATGGGATAAAAGATA
AAAAATTCATGCTGCACTACGAGTTTCTCCTTATGCAACTAATGAAATGGCAAAGTCACTGGTTTAAATAGAA
GAGAACTTGGGCATGGTGCTCTTGCTGAGAAAGCTTTGTATCCTGTTATTTCCCGGAGATTTTCTTTTACCATAA
GAGTTACATCTGAAGTCCTAGAGTCAAAATGGGTCTCTTCTATGGCATCTGCATGTGGCGGAAGTTTAGCATTA
TGGATTCAAGGGTTCCAATTTCTCTGCTGTTGCAGGCGTAGCAATAGGATTGGTCACCAAAACCGATCCTGAGA
AGGGTGAAATAGAAGATTATCGTTTGCTGACAGATATTTTGGGAATTGAAGATTACAATGGTGACATGGACTTCA
AAATAGCTGGCACTAATAAAGGAATAACTGCATTACAGGCTGATATTAATTAACCTGGAATACCAATAAAAATTG
TGATGGAGGCTATTCAACAAGCTTCAGTGGCAAAAAGGAGATATTACAGATCATGAACAAACTATTTCAAAC
CTCGAGCATCTAGAAAAGAAAATGGACCTGTTGTAGAACTGTTTCAAGTTCCATTATCAAAACGAGCAAAATTTG
TTGGACCTGGTGGCTATAACTTAAAAAACTTCAGGCTGAAACAGGTGTAACCTATTAGTCAGGTGGATGAAGAAA
CGTTTTCTGTATTGTCACCAACACCCAGTGCTATGCATGAGGCAAGAGACTTCATTACTGAAATCTGCAAGGATG
ATCAGGAGCAGCAATTAGAATTTGGAGCAGTATATACCGCCACAATAACTGAAATCAGAGATACTGGTGTAAATGG
TAAAATTATATCCAAATATGACTGCGGTACTGCTTCATAACACACAACCTTGATCAACGAAAGATTAAACATCCTA
CTGCCCTAGGATTAGAAGTTGGCCAAGAAATTCAGGTGAAATACTTTGGACGTGACCCAGCCGATGGAAGAATGA
GGCTTCTCGAAAAGTGCTTCAGTCGCCAGCTACAACCGTGGTCAGAACTTTGAATGACAGAAGTAGTATTGTAA
TGGGAGAACCTATTTACAGTCATCATCTAATTCAGTGATTTTTTTTTTAAAGAGAATTCTAGAATTCTATT
TTGTCTAGGGTGATGTGCTGTAGAGCAACATTTTAGTAGTATCTTCCATTGTGTAGATTTCTATATAATATAAAT
ACATTTTAATTATTTGTACTAAAATGCTCATTACATGTGCCATTTTTTAAATTCGAGTAACCCATATTTGTTTA
ATTGTATTTACATTATAAATCAAGAAATATTTATTATTAAGTAAGTCATTATACATCTTAGAAAAAATTACA
TAGTTTTGTTTTTACAATCTGAAATATATGAAAACTTAGATAGAATATGTCATATGTTATTATAACAGATCTC
TTCTACATCTTATTTTCTTCTTTGTATACAGTAGTGACAGTTTACCTTCAGTCATCCTACAGGTCACCTCTGT
GCCCAACAACCTTTACTTGCCCTACAACGGTTCACAACTGGAACAGGCCATATTCAAAGCCAGTGCCATTTCTTC
AGAACTGTTAACAGATATAGTGAGTTGAGGGAGCTAATCTGATACACTTTTGATAATATAATGCCTTTCAAATTA
GTTACCAAATCATAAACAGAGTGGAATAAATAAATGAGATTCTAACTAGGATGAATGTGGTAGTAAATGATGTA
TATTTTCAAATTTACCTAAACAGAAAGATTATTGGCTAAGGCAGGCGGATCATGAGATCAGGAGATCGAGACCA
TCCTGGCTAACACGGTGAAACCCCGTCTCTACTAAAAGTGCAAAAATTAGCCGGCGGTGGTGGCGCATGCCTGTA
ATCCTAGCTACTCAGGAGGCTGAGGCAGAAGAATCACTTCAACCCAGGAGGTGGAGGTTGTAGTGAGCCGAGATT
CGCCATTGCACTCCAGCTCAGGCAACAAGAGCAAACTCCGTCTC

810/6881

FIGURE 754

MAACRYCCSCLRLRPLSDGPFLLPRRDRAITQLQVRALWSSAGSRAVAVDLGNRKLEISSGKLARFADGSAAVQS
GDTAVMVTAVSKTKPSQFMPLVVDYRQKAAAAGRIPTNYLRREIGTSDKEILTSRIIDRSIRPLFPAGYFYDT
QVLCNLLAVDGVNEPDVLAINGASVALSLSDIPWNGPVGAVRIGIIDGEYVVPTRKEMSSSTLNLVVAGAPKSQ
IVMLEASAENILQQDFCHAIKVGVKYTQQIIQGIQQLVKEITGVTKRTPQKLFPSPEIVKYTHKLAMERLYAVFT
DYEHDKVSRAEAVNKIRLDTEEQLKEKFPEADPYEIIIESFNVAKEVFRSIVLNEYKRCRDLTSLRNVSCVD
MFKTLHGSALFQRGQTQVLCTVTFTDSLESIGKSDQVITAINGIKDKNFMLHYEFPPYATNEIGKVTGLNRRELGH
GALAEKALYPVIPRDFPFTIRVTSEVLESNGSSSMASACGGSLALMDSGVPISAVAGVAIGLVTKTDPEKGEIE
DYRLTDLGIEDYNGDMDFKIAGTNKGITLQADIKLPGIPKIVMEAIQQASVAKKEILQIMNKTISKPRASR
KENGPPVETVQVPLSKRAKFGPGGYNLKKLQAETGVTISQVDEETFVVFAPTPSAMHEARDFITEICKDDQEQQ
LEFGAVYTATITEIRDTGVMVKLYPNMTAVLLHNTQLDQRIKHPTALGLEVGQEIQVKYFGRDPADGRMRLSRK
VLQSPATTVVRTLNDRSSIVMGEPISQSSSNSQ

811/6881
FIGURE 755

GACAAGATGGCCACACCGGCGGTACCAGCAAGTGCTCCTCCGGCCACGCCAGCCCCAGTCCCGGCGGCGGTCCCA
GCCTCTGCCCCAGCCTCAGTTCCAGCGCCAACGCCAGCACCGGCTGCGGCTCCGGTTCCCGCTGCGGCTCCAGCC
TCATCCTCAGACCCTGCGGCAGCATCGGCTACAAC TGCGGCTCCTGGCCAGACCCCGGCCTCAGCGCAAGCTCCA
GCGCAGACCCAGCGCCCGCTCTGCCTGGTCCTGCTCTTCCAGGGCCCTTCCCCGGGCGCGCGTGGTCAGGCTG
CACCCAGTCATTTTGGCCTCCATTGTGGACAGCTACGAGAGACGCAACGAGGGTGCTGCCCCGAGTTATCGGGACC
CTGTTGGGAAGTGTGACAAACACTCAGTGGAGGTCACCAATGCTTTTCAGTGCCGCACAATGAGTCAGAAGAT
GAAGTGGCTGTTGACATGGAATTTGCTAAGAATATGTATGAAGTGCATAAAAAAGTTTCTCCAAATGAGCTCATC
CTGGGCTGGTACGCTACAGGCCATGACATCACAGAGCACTCTGTGCTGATCCATGAGTACTACAGCCGAGAGGCC
CCCAACCCCATCCACCTCACTGTGGACACAAGTCTCCAGAACGGCCGCATGAGCATCAAAGCCTATGTCAGCACT
TTAATGGGTGTCCCTGGGAGGACCATGGGAGTGATGTTTACACCTCTGACAGTGAAATACGCATACCTATGACACT
GAACGCATCGGAGTTGAGCTGATCATGAAGACCTGCTTTAGCCCCAACAGAGTGATTGGACTCTTAAGTGACTTG
CAGCAAGTAGGAGGGGCATCAGCTCGCATCCAGGATGCCCTGAGTACAGTGTTGCAATATGCAGAGGATGTACTG
TCTGGAAAGGTGTCAGCTGACAATACTGTGGGCCGCTTCCTGATGAGCCTGGTTAACCGAGTACCGAAAAATAGTT
CCCGATGACTTTGAGACCATGCTCAACAGCAACATCAATGACCTTTTGATGGTGACCTACCTGGCCAACCTCACA
CAGTCACAGATTGCCCTCAATGAAAACTTGTAACCTGTGAATGGACCCCAAGCAGTACACTTGCTGGTCTAGG
TATTAACCCAGGACTCAGAAGTGAAGGAGAAATGGGTTTTTTGTGGTCTTGAGTCACACTGAGATAGTCAGTTG
TGTTGACTCTAATAAACGGAGCCTACCTTTTGT

812/6881

FIGURE 756

MATPAVPASAPPATPAPVPAAVPASAPASVPAPTPAPAAAPVPAAPASSSDPAAASATTAAPGQTPASQAQAPAQ
TPAPALPGPALPGPFPGGRVVRLHPVILASIVDSYERRNEGAARVIGTLLGTVDKHSVEVTNCFVPHNESEDEV
AVDMEFAKNMYELHKKVSPNELILGWYATGHDITEHSVLIHEYYSREAPNPIHLTVDTSLQNGRMSIKAYVSTLM
GVPGRIMGVMFTPLTVKYÄYYDTERIGVELIMKTCFSPNRVIGLLSDLQQVGGASARIQDALSTVLQYAEDVLSG
KVSADNTVGRFLMSLVNRVPKIVPDDFETMLNSNINDLLMVTYLANLTQSQIALNEKLVNL

813/6881
FIGURE 757

GTTGGAGACTCGATTGTTGATGACAGCGGAAGAATGATAACAAAATGCGCGAGCGAGATAGTGAGCCGTTCTCCA
ACCCTTTGGCCCCCTGATGGCCACGATGTGGATGATCCTCACTCCTTCCACCAATCAAACTCACCAATGAAGACT
TCAGGAAACTTCTCATGACCCCCAGGGCTGCACCTACCTCTGCACCACCTTCTAAGTCACGTCACCATGAGATGC
CAAGGGAGTACAATGAGGATGAAGACCCAGCTGCACGAAGGAGGAAAAAGAAAAGTTATTATGCTAAGCTACGCC
AACAAGAAATTGAGAGAGAGAGAGAGCTAGCAGAGAAGTACCGGGGTCGTGCCAAGGAACGGAGAGATGGAGTGA
ACAAAGATTATGAAGAAACCGAGCTTATCAACACCACAGCTAACTACAGGGCTGTTGGCCCCACTACTAAGGCGG
ACAAATCAGCTGCAGAGAAGAGAAGACAGTTGATCCAGGAGTCCAAATTCTTGGGTGGTGACATGGAACACACCC
ATTTGGTGAAAGGCTTGGATTTTGTCTGTCTTCAAAAGGTACGAGCTGAGATTGCCAGCAAAGAGAAAGAGGAAG
AGGAACTGATGGAAGGCCCGAGAAAGAAACCAAGAAAGATGAGGATCCTGAAAATAAAATTGAATTTAAACAC
GTCTGGGCCGCAATGTTTACCGAGTGCTTTTAAGAACAAGCATATAAGCGGAATGAGTTGTTCTCGCCGGGCCG
CATGGCCTATGTGGTAGGCCTGGATGATGAGTAAGCTGACACAGATATCCCCACCCTCTTATCCGCAGCAAGGT
TGACTGCCCCACCATGGAGGCCCGAGACCATACTGACCACAAATGATATTGTCATTAGCAAGCTTACCCAGTTGCT
TTCATACCTGAGGCAGGGAACCCGTAACAAGAAGCTTAAGAAGAAGGATAAAGGGAAGCTGGAAGAGAAGAAACC
TGCTGAGGCTGACAGGAGTATTTTCGAAGACATTGGGGATTATACACCCTCCACAATCAAGACACCTTGGGACAA
GGAGCGGGAGAGATATCGGGAACGGGAGCGTGATCAGGAGACAGAGACCGTGACCGAGAGCGAGAGCGAGAACGA
GATCAGGAGCGAGAACGAGATCGGGAACGAGAGAGAGAAGAGGAAAAGAAGAGACACAGCTACTTTGAGAAGCCA
AAAGTACATGATGAGTCCGTGGACGTTGACAAAGGACCTGGGTCGGCCAAGGAGTTGATCAAGTCCATCAATGAA
AAGTTTGTGGGCTGTGCTGGCTGGGAAGGCACAGAATCGCTGAAAATTGCAGAAGACAAAAGCAGCTGAGAGAT
TTCTTTGGCATGTCCAACAGTTATGCAGAGTGCTATCCAGCCACGATGGATGACATGGCTGTGGATAGTGATGAG
GAGGTGGATTATAGCAAAATGGACCAGGGTAACAAGAAGAGAACCTTAAGCCGTTGGGACTTTGATACCCAGGAA
GACTACAGCGAGTATATGAACAACAAAGAGGCTTTGCCAAGGCTGCATTCCAGTATGGTATCAAAATGTCTGAA
GGGCGGAAAACAGACGCTTCAAGGAAACCAATGACAAAGCAGCGCTCGATTGCCAGTGAAGATTAGTGCAATC
ATTGAGAAGAGGAAGAAGATGGAAGCTGATGGGGTTGAAGTCAAAAGACCAAAATACTAA

814/6881

FIGURE 758

MPERDSEPF SNPLAPDGHVDVDPHSFHQSKLTNEDFRKLLMTPRAAPTSAPPSKSRHHEMPREYNEDEDPAARRR
KKKSYYAKLRQQEIERERELAEKYRGRAKERRDGVNKDYEETELINTTANYRAVGPTTKADKSAAEKRRQLIQES
KFLGGDMEHTHLVKGLDFALLQKVRAEIASKEKEEEEELMEKPQETKKDEDPENKIEFKTRLGRNVYRVLFKNKH
ISGMSCSCRAAWPMW

815/6881

FIGURE 759

ATGCCCCGAGAATGTGGCACCCCGAGCGGGGCGACTGCCGGGGCTGCCGGCGGCCGCGGGAAAGGCGCCTATCAG
GACCGCGACAAGCCAGCCCAGATCCGCTTCAGCAACATTTCCGCCGCCAAAGCGGTTGCTGATGCTATTAGAACA
AGCCTTGGACCAAAAGGAATGGATAAAATGATTCAAGATGGAAGGTGATGTAACCATTACAAATGATGGTGCT
ACCATTCTGAAACAAATGCAAGTATTACATCCAGCAGCCAGAATGCTGGTGGAGCTGCTAAGGCTCAAGATATA
GAAGCAGGAGATGGCACCACATCAGTAGTCATCATTGCTGGCTCCCTCTTAGATTCTTGTACCAAGCTTCTTCAG
AAAGGGATTCAATCAACCATCATTTCTGAGTCATTCCAGAAGGCCCTGGAAAAGGGCATTGAAATCTTGACTGAC
ATGTCTCGACCTGTGGAAGTGAAGTACAGAGAACTTTGTTAAATAGTGCAACCACTTCACTGAACTCAAAGGTG
GTTTCTCAGTATTCAAGTCTGCTTTCTCCAATGAGTGTAATGCAGTGATGAAAGTGATTGACCCAGCCACAGCC
ACCAGTGTAGATCTTAGAGATATTAATAAGTGAAGCTTGGTGGGACAATTGATGACTGTGAGTTGGTGGAA
GGGCTGGTTCTACCCAAAAAGTGTCAAATCTGGCATAACCAGAGTTGAAAAGGCCAAGATTGGGCTTATTTCAG
TTTTGCTTATCTGCTCCCAAAACAGACATGGATAATCAAATAGTGGTTCTGACTATGCCAGATGGACCGAGTG
CTGCGAGAAGAGAGAGCCTATATTTTAAATTTAGTGAAGCAAATTAACAAAAACAGGATGTAATGTCCTTCTCATA
CAGAAATCTATTCTAAGAGATGCTCTTAGTGATCTTGCAATTACACTTTCTGAATAAAATGAAGATCATGGTGATT
AAGGATATTGAAAGAGAAGACATTGAATTCATTTGTAAGACAATTGGAACCAAGCCAGTTGCTCATATTGACCAA
TTTACTGCTGACATGCTGGGTTCTGCTGAGTTAGCTGAGGAGGTCAATTTAAATGGTTCTGGCAAACCTGCTCAAG
ATTACAGGCTGTGCCAGCCCTGGAAAAACAGTTACAATTGTTGTTTCGTGGTTCTAACAAACTGGTGATTGAAGAA
GCTGAGCGCTCCATTTCATGATGCCCTATGTGTTATTTCGTTGTTTAGTGAAGAAGAGGGCTCTTATTGCAGGAGGT
GGTGCTCCAGAAATAGAGTTGGCCCTAGCATTAACTGAATATTACGAACACTGAGTGGTATGGAATCCTACTGC
GTTTCGTGCTTTTGCAGATGCTATGGAGGTCAATTCATCTACACTAGCTGAAAATGCCGGCCTGAATCCCATTCT
ACAGTAACAGAACTAAGAAACCGGCATGCCCAGGGAGAAAAAACTGCAGGCATTAAATGTCCGAAAGGGTGGTATT
TCCAACATTTTGGAGGAAGTGGTTGTCCAGCCTCTGTTGGTATCAGTCAGTGCTCTGACTCTTGCAACTGAACT
GTTTCGGAGCATTCTGAAAATAGATGATGTGGTAACACTCGATAATCTGGATAACTGACTAGCACCATTATGATC
ACCAGTATTGTGGCTGGAATGGAAGAAGATCACCTGGTGTTCTTGGTTGGAAGATTATTCCTCTGAATTTCT
GGGCTTGGTCTTCCAGTTGGCATTTCCTGAAGTTGATTGAAACAATTTAATGAAAATATTAAATATTTGGTTT
CAAAAGGCAGATTATCTTCTCCCAACATTCTGTTATTCTGATACTTTTGAAAACTAATAAAAACTAATAAAA
GAAGCGTA

816/6881
FIGURE 760

MPENVAPRSGATAGAAGGRGKGAYQDRDKPAQIRFSNISAACAVADAIRTS LGPKGMDKMIQDGKGDVTITNDGA
TILKQMQLHPAARMLVELSKAQDIEAGDGTTSVVIAGSLLDSCTKLLQKGIHPTIISSEFQKALEKGIEILTD
MSRPVELSDRETLNLSATTSLNSKVVSQYSSLLSPMSVNAV MKVIDPATATSVDLRDIKIVKKLGGTIDDCELVE
GLVLTQKVSNSGITRVEKAKIGLIQFCLSAPKTDMDNQIVVSDYAQMDRVLREERAYILNLVKQIKKTGCNVLII
QKSILRDALSDALHFLNKMIMVIKDIEREDIEFICKTIGTKPVAHIDQFTADMLGSAELAEVNLNGSGKLLK
ITGCASPGKTVTIVVRGSNKLVEEAAERSIHDALCVIRCLVKKRALIAGGGAPEIELALALTEYSRTLSGMESYC
VRAFADAMEVIPSTLAENAGLNP ISTVTELRNRHAQGEKTAGINVRKGGISNILEELVVQPLLVSVSALT LATET
VRSILKIDDVVNT

817/6881
FIGURE 761A

AGGAAGGAAGGAGCAGTTGGTTCAATCTCTGGTAATCTATGCCAGCAATTATGACAATGTTAGCAGACCATGCGAG
CTCGTCAGCTGCTTGATTTTCAGCCAAAACTGGATATCAACTTATTAGATAATGTGGTGAATTGCTTATACCATG
GAGAAGGAGCCAGCAAGAATGGCTCAAGAAGTACTGACACATTTAAAGGAGCATCCTGATGCTTGGACAAGAG
TCGACACAATTTTGGAAATTTCTCAGAATATGAATACGAAATACATGGAATACAAATTTTGGAAAATGTGATAA
AAACAAGGTGGAAGATTCTTCCAAGGAACAGTGCAGGAAGGAATAAAAAATACGTTGTTGGCCTCATTATCAAGA
CGTCATCTGACCCAACTTGTGTAGAGAAAGAAAGGTGTATATCGGAAAATTAATATGATCCTTGTTCAGATAC
TGAAACAAGAATGGCCCAACATTGGCCAACTTTTATCAGTGATATTGTTGGAGCAAGTAGGACCAGCGAAAGTC
TCTGTCAAAATAATATGGTGATTCTTAAACTCTTGAGTGAAGAAGTATTGATTTCTCTAGTGACAGATAACCC
AAGTCAAACTAAGCATTATAAAGACAGCATGTGCAATGAATTCTCACAGATATTTCACTGTGTGAGTTTGTAA
TGGAAAATTTCTCAAAATGCTCCACTTGTACATGCAACCTTGGAAACATTGCTCAGATTTCTGAAGTGGATTCCCC
TGGGATATATTTTGGAGACCAATTAATCAGCACATTGATTTATAAGTTCCTGAATGTTCCAATGTTTCGAAATG
TCTCTCTGAAGTGCCCTCACTGAGATTGCTGGTGTGAGTGAAGCCAATATGAAGAACAATTTGTAACACTATTTA
CTCTGACAATGATGCAACTAAAGCAGATGCTTCCTTTAAATACCAATATTCGACTTGCGTACTCAAATGGAAAG
ATGATGAACAGAAGTTCAATTCAAATCTCAGTTTGTCTCTGCACCTTCTTAAGGAACATGATCAACTTATAG
AAAAAGATTAAATCTCAGGGAAGCTCTTATGGAGGCCCTTCATTATATGTTGTTGGTATCTGAAGTAGAAGAAA
CTGAAATCTTTAAATTTGTCTTGAATACTGGAATCATTGGCTGCTGAAGTCTATAGAGAGAGTCCATTCTCTA
CATCTGCCTCTCCGTTGCTTTCTGGAAGTCAACATTTTGATGTTCTCCAGGAGACAGCTATATTTGCCCATGT
TATTCAAGGTCCGTTTATTAATGGTTAGTGAATGGCTAAACCAGAGGAAGTATTGGTTGTAGAGAATGATCAAG
GAGAAGTTGTGAGAGAATTCATGAAGGATACAGATTCCATAAATTTGTATAAGAATATGAGGGAACATTTGTTT
ATCTTACTCATCTGGATTATGTAGATACAGAAAGAATAATGACAGAGAAGCTTCACAATCAAGTGAATGGTACAG
AGTGGTCATGGAATAATTTGAATACATTGTGTTGGGCAATAGGCTCCATTAGTGGAGCAATGCATGAAGAGGACG
AAAAACGATTTCTTGTACTGTTATAAAGGATCTATTAGGATTATGTGAACAGAAAAGAGGCAAGATAATAAAG
CTATTATTGCATCAAATATCATGTACATAGTAGGTCAATACCCACGTTTTTTGAGAGCTCACTGGAAATTTCTGA
AGACTGTAGTTAACAAGCTGTTTCAATTCATGCATGAGACCCATGATGGAGTCCAGGATATGGCTTGTGATACTT
TCATTAAATAGCCCAAAATGCCGCGAGGCATTTCTGTTGAGTTTCAAGTTGAGAGTATGCCATTTATTGATG
AAATTTGAACAACATTAACACTATTATTGTGATCTTCAGCCTCAACAGGTTTCATACGTTTTATGAAGCTGTGG
GGTACATGATTGGTGCAAAACAGATCAAACAGTACAAGAGCACTTGATAGAAAAGTACATGTTACTCCCTAATC
AAGTGTGGGATAGTATAATCCAGCAGGCAACCAAAATGTGGATATACTGAAAGATCCTGAAACAGTCAAGCAGC
TTGGTAGCATTTTGAACAAATGTGAGAGCCTGCAAGCTGTTGGACACCCCTTTGTAATTCAGCTTGGAAAGAA
TTTATTAGATATGCTTAATGTATACAAGTGCCTCAGTGAAATATTTCTGCAGCTATCCAAGCTAATGGTGAAA
TGGTTACAAAGCAACCATTGATTAGAAAGTATGCGCAACTGTAAAGGGAACTTTAAAGTTAATATCTGGTTGGG
TGAGCCGATCCAATGATCCACAGATGGTCGCTGAAAATTTTGTTCCTCTGTTGGATGCAGTTCTCATTGATT
ATCAGAGAAATGTCCAGCTGTAGAGAACCAGAAAGTGCTTAGTACTATGGCCATAATTGTCAACAGTGGGG
GACATATAACAGCTGAAATACCTCAAATATTGATGCTGTTTTTGAATGCACATTGAATATGATAAATAAGGACT
TTGAAGAATATCCTGAACATAGAACGAACCTTTTCTTACTACTTCAGGCTGTCAATTCTCATTGTTTCCAGCAT
TCCTTGCTATTCCACCTACACAGTTTTAACTTGTTTTGGATTCCATCATTGTTGGCTTTCAAACATACTATGAGGA
ATGTCGCAGATACGGGCTTACAGATACTTTTACACTCTTACAAAATGTTGCACAAGAAGAAGCTGCAGCTCAGA
GTTTTTATCAAACCTTATTTTGTGATATTCTCCAGCATATCTTTCTGTTGTGACAGACACTTCACATACTGCTG
GTTTAAACAATGCATGCATCAATTCTTGATATATGTTTAAATTTGGTTGAAGAAGGAAAAATAAGTACATCATTA
ATCCGTGGAATCCAGTTAACAACCAATCTTTCTTCAGGAATATGTGGCTAATCTCCTTAAGTCGGCCTTCCCTC
ACCTACAAGATGCTCAAGTAAAGCTCTTTGTGACAGGGCTTTTCAGCTTAAATCAAGATATTCCTGCTTTCAAGG
AACATTTAAGAGATTTCTAGTTCAAATAAAGGAATTTGCAGGTGAAGACACTTCTGATTTGTTTTTGGAAAGAGA
GAGAAATAGCCCTACGGCAGGCTGATGAAGAGAAACATAAACGTCAAATGTCTGCTCCCTGGCATCTTAAATCCAC
ATGAGATTCCAGAAGAAATGTGTGATTAAATCCAAATTCATGCTGTTTTTTTTCTCTGCAACTCCGTTAGCAGA
GGAAAACAGCATGTGGGTATTGTGCGACCAAAATGATGCCAATTTGTAAATTAATGTACCTAGTGGCCCTTT
TTCTTATGTGTTTTTTTGTATAAGAAATTTCTGTGAAATATCCTTCCATTGTTTAAAGCTTTTGTGTTTGGTCATC
TTTATTTAGTTTGCATGAAGTTGAAATTAAGGCATTTTAAAAATTTTACTTCATGCCCATTTTTGTGGCTGGG
CTGGGGGGAGGAGGCAATTCATTTGAACATATACTTGTAAATCTAATGCAAAATTATACAATTTTCTCTGTAA

818/6881

FIGURE 761B

ACAATACCAATTTTAAATTAGGGAGCATTTTCCTTCTAGTCTATTTTCAGCCTAGAAGAAAAGATAATGAGTAAAA
CAAATTGCGTTGTTTAAAGGATTATAGTGTGCTGCATTGTCTGAAGTTAGCACCTCTTGGACTGAATCGTTTGTCTA
GACTACATGTATTACAAAGTCTCTTTGGCAAGATTGCAGCAAGATCATGTGCATATCATCCCATTGTAAAGCGAC
TTCAAAAATATGGGAACACAGTTAGTTATTTTACACAGTTCCTTTTGTGTTTGTGTGTGTGTGCTGTCGCTTGT
CGACAACAGCTTTTGTGTTTCCTCAATGAGGAGTGTGCTCATTGTGAGCCTTCATTAACTCGAAGTGAAATGG
TTAAAAATATTTATCCTGTTAGAATAGGCTGCATCTTTTAACTCACTTAAAAAACAACAACTCTGGCTTT
TGAGATGACTTATACTAATTTACATTGTTTACCAAGCTGTAGTGCTTTAAGAACACTACTTAAAAAGCAAAATAA
ACTTGTTTACATTTAAAAAAA

819/6881
FIGURE 762

MPAINTMLADHAARQLLDIFSQKLDINLLDNVNCCLYHGEGAQORMAQEVLTILKEHPDAWTRVDTILEFSQNMNT
KYYGLQILENVIKTRWKILPRNQCEGIKKYVVGLIIKTSSDPTCVEKEKVYIGKLNMLVQILKQEWPKHWPTFI
SDIVGASRTSESLCQNNMVLKLLSEEVDFSSGQITQVSKHLKDSMCNEFSQIFQLCQFVMENSQNAPLVHAT
LETLLRFLNWIPLYIFETKLITLIYKFLNVPFRNVSLKCLTEIAGVSVSQYEEQFVTLFTLTMMQLKQMLPL
NTNIRLAYSNGKDDEQNFIQNLSLFLCTFLKEHDQLIEKRLNLRETLMELHYMLLVSEVEETEIFKICLEYWNH
LAAELYRESPFSTSASPLLGSQHFDVPPRRQLYLPMLFKVRLLMVSRMAKPEEVLVVENDQGEVVREFMKDTS
INLYKNMRETLVYLTHLDYVDTERIMTEKLHNQVNGTEWSWKNLNTLCWAIGSISGAMHEEDEKRFLVTVIKDLL
GLCEQKRGKDNKAIIASNIMYIVGQYPRFLRAHWKFLKTVVNKLFEFMHETHDGVQDMACDTFIKIAQKCRHFV
QVQVGEVMPFIDEILNNINTIICDLQPQVHTFYEAAGYMIGAQTDTQTVQEHLEKYMLLPNQVWDSIIQQATKN
VDILKDPETVKQLGSILKTNVRACKAVGHFVIQLGRIYLDMLNVYKCLSENISAAIQANGEMVTKQPLIRSMRT
VKRETLKLISGWVSRSDNPQMVAENFVPPLLDAVLIDYQRNVAAREPEVLSTMAIIVNKLGGHITAEIPQIFDA
VFECTLNMINKDFEYPEHRTNFFLLQAVNSHCFFAFLAIPPTQFKLVLDSEIWAFAKHTMRNVADTGLQILFTL
LQNVAQEEAAAQSFYQTYFCDILQHIFSVVTDTSHTAGLTMHASILAYMFNLVEEGKISTSLNPGNPVNNQIFLQ
EYVANLLKSAPPHLQDAQVKLFVTGLFSLNQDIPAFKEHLRDFLVQIKEFAGEDTSDLFLEEREIALRQADEEKH
KRQMSVPGIFNPHEIPEEMCD

820/6881
FIGURE 763

GGCGGCGGCAGCAGCTGCTTGGGCGCCGTGCGGTGGTGA CTGAGCTACGAGCCTGGCAGCAGGTGTGCGCCGAGC
CCCGGCCTGGCCCGCCCCCGCGTGCTCC CAGGCTCCGCACCCCTGATGCTGCGCGGGTGCTGAGCCCACTTCG
GCCAGGACGATGGTTAAGTATTTCTGGGCCAGAGCGTGCTCCGGAGTTCTGGGACCAAGTGTTGCGCCGCCTTC
TGGCAGCGGTACCCGAATCCCTATAGCAAACATGTCTTGACGGAAGACATAGTACACCGGGAGGTGACCCCTGAC
CAGAAACTGCTGTCCCGGCGACTCCTGACCAAGACCAACAGAATGCCACGCTGGGCCAAGCAACTATTTCTGCC
AATGTTGCTCACTCGGTGTACATCCTGGAGGACTCTATTGTGGACCCACAGAATCAGACCATGACTACCTTCACC
TGGAACATCAACCACGCCCCGGCTGATGGTGGTGGAGGAACGATGTGTTTACTGTGTGA ACTCTGACAACAGCGGC
TGGACTGAAATCCGCCGGGAAGCCTGGGTCTCCTCTAGCTTATTTGGTGTCTCCAGAGCTGTCCAGGAATTTGGT
CTTGCCCAGTTCAA AAGCAACGTGACCAAGACTATGAAGGGTTTTGAATATATCTTGACTAAGCTGCAAGGCGAG
GCCCCTTCCAAAACACTTGAGACAGCCAAGGAAGCCAAGGAGAAGGCAGAGGAGACGGCACTGGCAGCTACAGAG
AAGGCCAAGGACCTCGCCAGCAAGGCGGCCACCAAGAAGCAGCAGCAGCAACAGTTTGTGTAGCCAGCCCAC
CACCACCACAGCAACCCAGACAGCTAGGCTTAGCCCCCTCTGCCCTCCCTCCATTGTACTTTATCATTAAAAATCA
ACTTCCA

821/6881
FIGURE 764

CTGACTCTCTGAGGCTCATTTTGCAGTTGTTGAAATTGTCCCGCAGTTTTCAATCATGTCTGAACCAATCAGAG
TCCTTGTGACTGGAGCAGCTGGTCAAATTGCATATTTACTGCTGTACAGTATTGGAAATGGATCTGTCTTTGGTA
AAGATCAGCCTATAATTCTTGTGCTGTTGGATATCACCCCATGATGGGTGTCCTGGACGGTGTCTAATGGAAC
TGCAAGACTGTGCCCTTCCCCTCCTGAAAGATGTCATCGCAACAGATAAAGAAGACGTTGCCTTCAAAGACCTGG
ATGTGGCCATTCTTGTGGGCTCCATGCCAAGAAGGGAAGGCATGGAGAGAAAAGATTTACTGAAAGCAAATGTGA
AAATCTTCAAATCCCAGGGTGCAGCCTTAGATAAAATACGCCAAGAAGTCAGTTAAGGTTATTGTTGTGGGTAATC
CAGCCAATACCAACTGCCTGACTGCTTCCAAGTCAGCTCCATCCATCCCCAAGGAGAACTTCAGTTGCTTGACTC
GTTTGGATCACAACCGAGCTAAAGCTCAAATTGCTCTTAAACTTGGTGTGACTGCTAATGATGTAAAGAATGTCA
TTATCTGGGGAAACCATTCTCGACTCAGTATCCAGATGTCAACCATGCCAAGGTGAAATTGCAAGGAAAGGAAG
TTGGTGTATTATGAAGCTCTGAAAGATGACAGCTGGCTCAAGGGAGAATTTGTCACGACTGTGCAGCAGCGTGGCG
CTGCTGTCATCAAGGCTCGAAAACCTATCCAGTGCCATGTCTGCTGCAAAAGCCATCTGTGACCACGTCAGGGACA
TCTGTTTGGAAACCCAGAGGGAGAGTTTGTGTCCATGGGTGTTATCTCTGATGGCAACTCCTATGGTGTTCTTG
ATGATCTGCTCTACTCATTCCCTGTTGTAATCAAGAATAAGACCTGGAAGTTTGTGGAAGGTCTCCCTATTAATG
ATTTCTCACGTGAGAAGATGGATCTTACTGCAAAGGAACTGACAGAAGAAAAAGAAAGTGCTTTTGAATTTCTTT
CCTCTGCCTGACTAGACAATGATGTTACTAAATGCTTCAAAGCTGAAGAATCTAAATGTCGTCTTTGACTCAAGT
ACCAATAATAATAATGCTATACTTAAATTACTTGTGAAAAACAACACATTTTAAAGATTACGTGCTTCTTGGTA
CAGGTTTGTGAATGACAGTTTATCGTCATGCTGTTAGTGTGCATTCTAAATAAATATATATTCAAATG

822/6881
FIGURE 765

MSEPIRVLTGAAGQIAYSLLYSIGNGSVFGKDQPIILVLLDITPMMGVLDGVLMELODCALPLLKDVIATDKED
VAFKDLDVAILVGSMRREGMERKDLLKANVKIFKSQGAALDKYAKKSVKVIVVGNPANTNCLTASKSAPSIPKE
NFSCLTRLDHNRKAQIALKLGVLTANDVKNVIIWGNHSSQYPDVNHAKVKLQKKEVGVEALKDDSWLKGEFVT
TVQQRGA AVIKARKLSSAMSAKAICDHVRDIWFGTPEGEFVSMGVISDGNSYGVPPDLLYSFPVVIKNKTWK FV
EGLPINDFSREKMDLTAKELTEEKESAFEFLSSA

823/6881

FIGURE 766

TCTTTTAAGTATACTCTCTACTTATTAAAAAAGTCCTTTCCACGCTACCTGCAGAGGGGTCCATACG
GCGTTGTTCTGGATTCCCATCGTAACTTAAAGGGAAACTTTCACAATGTCCGGAGCCCTTGATGTCCTGCAAATG
AAGGAGGAGGATGTCCTTAAGTTCCTTGCAGCAGGAACCTCACTTAGGTGGCACC AATCTTGACTTCCAGATGGAA
CAGTACATCTATAAAAGGAAAAGTGATGGCATCTATATCATAAATCTGAAGAGGACCTGGGAGAAGCTTCTGCTG
GCAGCTCGTGCTATTGTTGCCATTGAAAACCTTGCTGATGTCAGTGTTATATCCTCCAGGAATACTGGCCAGAGG
GCCGTGCTAAAGTTTGCTGCTGCCACTGGAGCCACTCCAATTGCTGGCCGCTTCACTCCTGGAACCTTCGCTAAC
CAGATCCAGGCAGCCTTCCGGGAGCCACGGCTTCTTGTTGTTACTGACCCAGGGCTGACCACCAGCCTCTCATG
GAGGCATCTTATGTTAACCTACCTACCATTGCGCTGTGTAACACAGATTCTCCTCTGCACTATGTGGACATTGCC
ATCCCATGCAACAACAAGGGAGCTCACTCAGTGGGTTTGATGTGGTGGATGCTGGCTCGGGAAGTTCTGCGCATG
CGTGGCACCATTTCCTGTGAACACCCATGGGAGGTCATGCCTGATCTGTACTTCTACAGAGATCCTGAAGAGATT
GAAAGAGAAGACAGGCTGCTGCTGAAAAGGCAGTGACCAAGGAGGAATTCAGGGTGAATGGACTGCTCCAGCT
CCTGAGTTCACTGTTACTCAGCCTGAGGTTGCAGACTGGTCTGAAGGTGTACAGGTGCCCTCTGTGCCATTCCC
TACTGAAGACGGGAGCGCTCAGCCTGCCACGGAAGACTGGTCTGCAGCTCCCACTGCTCAGGCCACTGAATGGGT
AGGAGCAACCACTGACTGGTCTTAAGCTGTTCTTG CATAGGCTCTTAAGCAACATGGAAAAATGGTTGATGGAAA
ATAAACATCAGTTTCT

824/6881
FIGURE 767

MSGALDVLQMKEEDVLKFLAAGTHLGGTNLDFQMEQYIYKRKSDGIYIINLKRTWEKLLLAARAIVAIENPADVS
VISSRNTGQRAVLKFAAATGATPIAGRFTPGTFANQIQAAFREPRLLVVTDPRAHQPLMEASYVNLPTIALCNT
DSPLHYVDIAIPCNNKGAHSVGLMWWMLAREVLRMRGTISCEHPWEVMPDLYFYRDP EEIEREEQAAAEKAVTKE
EFQGEWTAPAPEFTVTQPEVADWSEGVQVPSVPIPY

825/6881

FIGURE 768

GCAGAGGGGTCCATACGGCGTTGTTCTGGATTCCCATCGTAACTTAAAGGGAACTTTCACAATGTCGGGAGCCC
TTGATGTCCTGCAAATGAAGGAGGAGGATGTCCTTAAGTTCCTTGACGACGGAACCTCACTTAGGTGGCACCAATC
TTGACTTCCAGATGGAACAGTACATCTATAAAAGGAAAAGTGATGGCATCTATATCATAAATCTGAAGAGGACCT
GGGAGAAGCTTCTGGCCGCTTCACTCCTGGAACCTTCGCTAACCAGATCCAGGCAGCCTTCCGGGAGCCACGGCT
TCTTGTGGTTACTGACCCCAGGGCTGACCACCAGCCTCTCATGGAGGCATCTTATGTTAACCTACCTACCATTGC
GCTGTGTAACACAGATTCTCCTCTGCACTATGTGGACATTGCCATCCCATGCAACAACAAGGGAGCTCACTCAGT
GGGTTTGATGTGGTGGATGCTGGCTCGGGAAGTTCTGCGCATGCGTGGCACCATTTCCTGTGAACACCCATGGGA
GGTCATGCCTGATCTGTACTTCTACAGAGATCCTGAAGAGATTGAAAGAGAAGAGCAGGCTGCTGCTGAAAAGGC
AGTGACCAAGGAGGAATTTTCAGGGTGAATGGACTGCTCCAGCTCCTGAGTTCACTGTTACTCAGCCTGAGGTTGC
AGACTGGTCTGAAGGTGTACAGGTGCCCTCTGTGC

826/6881
FIGURE 769

MSGALDVLQMKEEDVLKFLAAGTHLGSTNLDFQMEQYIYKRKSDGIYIINLKRTWEKLLASLLEPSLTRSRQPSG
SHGFLWLLIPGLTTSLSQRHLKLTYP LLCVTQILLCAMWTLPSHATARELTQWV ▾

827/6881
FIGURE 770

GGCCCAGCTGCTGAGAGGAGTTGCCTGAGAGTGACCTTTGCATCTGCCTGTCCAGCCAGCATGGAACCAAAGCGG
ATCAGAGAGGGCTACCTTGTGAAGAAGGGGAGCGTGTTCAATACGTGGAAACCCATGTGGGTTGTATTGTTAGAA
GATGGAATTGAATTCATATAAGAAGAAAAGTGACAACAGCCCCAAAGGAATGATCCCCTGAAAGGGAGCACTCTG
ACTAGCCCTTGTCAAGACTTTGGCAAAAGGATGTTTGTGTTTAAGATCACTACGACCAAACAGCAGGACCCTTC
TTCCAGGCAGCCTTCTTGGAGGAGAGAGATGCCTGGGTTTCGGGATATCAAGAAGGCCATTAAATGCATTGAAGGA
GGCCAGAAATTTGCCAGGAAATCTACCAGGAGGTCCATTGACTGCCAGAAACCATTGACTTAGGTGCCTTATAT
TTGTCCATGAAAGACACTGAAAAAGGAATAAAAGAACTGAATCTAGAGAAGGACAAGAAGATTTTAAATCACTGC
TTCACAGGTAACCTGCCTCATTGATTGGCTGGTATCCAACAGTCTGTTAGGAATCGCCAGGAAGGCCCTCATGATT
GCTTCATCGCTGCTCAATGAGGGGTATCTGCAGCCTGCTGGAGACATGTCCAAGAGTGCAGTGGATGGAACCTGCT
GAAAACCCCTTTCCTGGACAACCTGATGCCCTTCTACTACTTTCCAGACAGTGGGTTCTTCTGTGAAGAGAATTCC
AGTGATGATGATGTGATTCTGAAAGAAGAAATTCAGAGGGGTTCATTATCAAGCAGGGATGTTTACTGAAGCAGGGG
CATAGAAGGAAAACTGAAAAGTGAGGAAGTTCATCTTGAGAGAAGACCTGCCTACCTGCACTACTATGACCCT
GCTGGGGCAGAAGATCCCTGGGAGCAATTCACTTGAGAGGCTGTGTGGTGACTTCAGTGGAGAGCAACTCAAAT
GGCAGGAAGAGTGAGGAAGAGAACCTTTTTGAGATCATCACAGCAGATGAAGTGCACTATTTCTTGAAGCAGCC
ACCCCAAGGAGCGCACAGAGTGGATCAGAGCCATCCAGATGGCCTCCCGAACTGGGAAGTAAAGAGACTCCTGC
ATTCTCTCTCCCTCTCTGAGGGAAGCCCATGGACAAGCTCAGTCCAGGACCTGTCCACTTCTGTGACAAATCAAC
GGGAAACAGCCCAGGGGTGGGAAGTTTTCAATTGAGGGGGGTCTGAATGTAACCTACCATGTGGTGTGCAAGGT
TCCCTGCAATTGTATTGCTCACTGCAGCCCTCTGCCCTATCCATGACCCCAAGCAGATATAACAAGCTGTGC
AGCCTCAGTAGGCTGCTTGCCCTCTCCAGGCCCTCAGGGCCTTCTTGAAAAATGAAGAAATTAACCTAGTAGATT
CCTGAGGTCCCCCTAGCTTAAAAAATAAATCTGCCCATGATTCTAACCTCGCAGTAGTGATAGTGATC
TAGTTGTTCTGCTGGTGTCTTCTTGGCTAAGTCTTGGCCTTCAGTTATCTTCAAATGTACCAGAACCTGAGCC
AACGCCTCCCTGTGAACTGTGTGCTGATCTGTAGTACAGTACCAGGAAGAAACCTCTTTTGTCTCTTTAGACAT
CTTCTACTTGTCTTGTGGCTTGAGATCGTGTAAACAAATGAAGGAGGGCTCTCTTCTTCTCTCATCCTACTC
AAAACTTCCCGAGAGCAGTGGTGGTTTTGAGGGTTTTGACTTCTATTACTTTTGGCAGCCTGGAAGTTGTGTC
TTCTGGGAAAGAGACCTGGGGAGGCCAGGAGTAGCTGAGGGTCTTCTGTGCCCTTAAACCGCCAGAGGAGCC
CTATTCCACTCTGGTTTTAGGCTGATCTGAGAGGGTCTCCCTTGTTCCTTTCTGGAGCATTTCTCTAACGTTTA
TTACAATTAGGAGGGGGACCCACATCTGTGAGATTCTGTTTCATTTGAGGTTTACAGAAAAAAGTGGCCA
GATGTGTTCCCCCATGGGTGAGAGGCCTGGGCACTGCCTGGTGAATGTGTCTTGGGCAGCTGCAGCAAGTGG
AGGGGCTGAACCTACTGGCCAGCTCACTGGATGATGGGTTAATACAACAACCTGCAGTGAAGGACTCAGAGCCACA
CAGAACTTCTGAGAGGGGCTGTAGCATTGCGCAGCATCTTCAGTTCTCCAGTAAATGATATTGCGTTCGTGCCT
CAGCTTTAAGCACAAGTAGCAGCAGCTCCTGCTTGAGTTCTGAGGGCATCATGGCCCTATGATTAAACAGAGTGA
TCTAACCTAGACTAAAATTGGGAACCTATTGCAATTTTGACCCTGACCACTAAGTAGTGATTCTTCTCCAAAA
TTGAGAAAGACAGCACCCATTGAAGCAGATATGTGTGTGAAGTATATTTTCAATCCAGATTTTAAATTTAA
GGCTCCAGGAAAGAAAGGAGAGTAGAACATTTTCTCATTTTATCAAATCCTCTCTTGCCCTCCCTCAATTCCC
CTGTAACATTCTGAAGCTGTCCCCTCCAGATGGTTTTATCAATAGCCTAGAGGTAAAGAACTGCTTTTTTC
TCTGATTCTTTAATAAATTATCTTTATAGAATATGCACAAGTTTTTCTACACTCAGTGTTAAAGTATTTATTAAT
GGGAAGTCAACTTAATGTTTTGAAATAAATATATGACTCTGTTTAAAT

828/6881
FIGURE 771

CTTTCGGGAGACTGGAGTCGAAGGCCGTGAGGTATTTTTCTAAGXXXXXXXXXXXXXXXXXXXXXXXXXXXXX
XXXXXXXXXAACCCCGGAGGAGGGTGAGGAGCAGAGCTGGCCATAATGGCAGGTGAAGAAATTAATGAAGACTAT
CCAGTAGAAATTCACGAGTATTTGTCAGCGTTTGAGAATTCATTGGTGCTGTGGATGAGATGCTGAAGACCATG
ATGTCTGTTTCTAGAAATGAGTTGTTGCAGAAGTTGGATCCACTTGAACAAGCAAAAGTGGATTTGGTTTCTGCA
TACACATTAAATTCAATGTTTTGGGTTTATTTGGCAACCCAAGGAGTTAATCCTAAGGAACATCCAGTAAACAG
GAATTGGAAAGAATCAGAGTATATATGAACAGAGTCAAGGAAATAACAGACAAGAAAAAGGCTGGCAAGCTGGAC
AGAGGTGCAGCTTCAAGATTTGTAAAAATGCCCTCTGGGAACCAAAATCGAAAAATGCATCAAAAGTTGCCAAT
AAAGGAAAAAGTAAAAGTTAACTTTTTGGTTTTGATGTACACATATTCAAAAAGTACATTAATATGTAAATTCAC
AGTAAATATGTAAAGCTAAATACTTTCTCTCCAAAGATCATTATCTTTATTGATTAGCACTGAGGATTTTAACA
TTGTGATATATTATATATTATAATTTACCATCTCTTGATGAGACTCTTATTTCTTTATATAGGTCAGTCTTGCA
AGTACCATTTTATAAGCAGCTGTGAAATTTAAGTGAAATGTTCTTTGTAAACATTTGTACTATTTTAAATGAATA
ATGACCTTATGAAGTATGCTATCTGTAGGCTGAAATTATAGGTACATCTGTTTTCACTATATGATATTAAGAAAG
CGTGAAATGACTTAAATGTTTCAATTTTTTCTGTATAGATACTTTATCATGTTTTCATGATTTTAGGAATTACTGC
TTTGTGATATTCAAAGTGTGAACTAAACTTTATGGTTGTACTTTAATTCTTGGCATGTTGCCTCTATGTCCC
ATTTAAATAAAATACATTCTCATTAACTTTAGATGGGAAATAAGGTTGTATGTTGATGGATGAATTTTGGCATG
ATGACTGTACTCTCAATAAAGGCTGAAAATGTTGT

829/6881

FIGURE 772

CCITTCACCTTCGCCCTCCAGCTGCTGCAGCTGCAGCCCCGACCGCGAGCGTGCCAAGCGGCTTCAGCAGCTAGCGGA
GCGGTGGCGGGCGGCCCCGCTCAGGAGACCACCAGATTCCCCCTCTTCCCGGGCCTCGCCATGGCGACCCACGGAC
AGACTTGCGCGCGTCCAATGTGTATTCTCCATCATATGCTGACCTTGGCAAAGCTGCCAGAGATATTTCAACA
AAGGATTTGGTTTTGGGTGGTGAAACTGGATGTGAAAACAAAGTCTTGCAGTGGCGTGGAATTTCAACGTCCG
GTTTCATCTAATACAGACACTGGTAAAGTTACTGGGACCTGGAGACCAAATACAAGTGGTGTGAGTATGGTCTGA
CTTTCACAGAAAAGTGAACACTGATAACACTCTGGGAACAGAAATCGCAATTGAAGACCAGATTTGTCAAGGTT
TGAAACTGACATTTGATACTACCTTCTCACCAAATCAAAGCTGACAAGGAATAACTTTGCAGTGGGCTACAGGAC
TGGGGACTTCCAGCTACACACTAATGTCAATGATGGGACAGAATTTGGAGGATCAATTTATCAGAAAGTTGTGA
AGATCTTGACACTTCAGTAAACCTTGCTTGGACATCAGGTACCAACTGCACTCGTTTTGGCATTGCAGCTAAATA
TCAGTTGGATCCCACTGCTTCCATTTCTGCAAAGTCAACAACCTCTAGCTTAATTGGAGTAGGCTATACTCAGAC
TC

830/6881
FIGURE 773

AGCTGCTGCAGCTGCAGCCCGACCGCGAGCGTGCCAAGCGGCTTCAGCAGCTAGCGGAGCGGTGGCGGCGGCCCC
GCTCAGGAGACCACCAGATTCCCCTCTTCCCGCGGCCCTGCCATGGCGACCCACGGACAGACTTGC GCGCGTCCA
ATGTGTATTCCCTCCATCATATGCTGACCTTGGCAAAGCTGCCAGAGATATTTTCAACAAAGGATTTGGTTTTGGG
TTGGTGAAACTGGATGTGAAAACAAAGTCTTGCA GTGGCGTGGAATTTTCAACGTCCGGTTCATCTAATACAGAC
ACTGGTAAAGTTACTGGGACCCTGGAGACCAAATACAAGTGGTGTGAGTATGGTCTGACTTTCACAGAAAAGTGG
AACACTGATAACACTCTGGGAACAGAAATCGCAATTGAAGACCAGATTTGTCAAGGTTTGAAACTGACATTTGAT
ACTACCTTCTCACCAAACACAGGAAAGAAAAGTGGTAAAATCAAGTCTTACAAGAGGGAGTGTATAAACCTTGGT
TGTGATGTTGACTTTGATTTTGTGGACCTGCAATCCATGGTTCAGCTGTCTTTGGTTATGAGGGCTGGCTTGCT
GGCTACCAGATGACCTTTGACAGTGCCAAATCAAAGCTGACAAGGAATAACTTTGCAGTGGGCTACAGGACTGGG
GACTTCCAGCTACACACTAATGTCAATGATGGGACAGAAATTTGGAGGATCAATTTATCAGAAAGTTTGTGAAGAT
CTTGACACTTCAGTAAACCTTGCTTGGACATCAGGTACCAACTGCACCTCGTTTGGCATTGCAGCTAAATATCAG
TTGGATCCCACTGCTTCCATTTCTGCAAAAGTCAACAACCTCTAGCTTAATTGGAGTAGGCTATACTCAGACTCTG
AGGCCTGGTGTGAAGCTTACACTCTCTGCTCTGGTAGACGGGAAGAGCATTAAATGCTGGAGGCCACAAGGTTGGG
CTCGCCCTGGAGTTGGAGGCTTAATCCAGCTGAAAGAAACCTTTGGGAATGGATATCAGAAGATTTGGCCTTAAT
ATATTTCCATTGTGACCAGCAGCAGGCTTTTTTCCCCCAAGAAGATGATCAAAACAAAGGATGATCTCAACAAGA
GCTGTATTTTAAGTATTTAGACAGTTCCTTTGTAGCTGGTTTCTAGTTGAATTGGTTATCTAGTTACCAATGCTG
CAGTCCTGCAGTCACCTATACATTATTTAAATGTATTTAACTGTTAAATGCGCTACCCACCAATAATGAAATAGA
CCTTTATGAAAA

831/6881
FIGURE 774

MATHGQTCARPMCIPPSYADLGKAARDIFNKGFGFGLVKLDVKTKSCSGVEFSTSGSSNTDTGKVTGTLETXYKW
CEYGLTFTEKWNTDNTLGTEIAIEDQICQGLKLTFTTSPNTGKKSGKIKSYKRECINLGCDDVDFDFAGPAIHG
SAVFGYEGWLAGYQMTFDSAKSLTRNNFAVGVRTGDFQLHTNVNDGTEFGGSIYQKVCEDLDTSVNLAWTSGTN
CTRFGIAAKYQLDPTASISAKVNNSSLIGVGYTQTLRPGVKLTLSALVDGKSINAGGHHKVGLEALELEA

832/6881
FIGURE 775

ATGGTCTGACTTTCACAGAAAAGTGGAACTGATAACACTCTGGGAACAGAAATCGCAATTGAAGACCAGATT
GTCAAGGTTTGAAACTGACATTTGATACTACCTTCTCACCAAACACAGGAAAGAAAAGTGGTAAAATCAAGTCTT
ACAAGAGGGAGTGTATAAACCTTGTTGTGATGTTGACTTTGATTTTGCTGGACCTGCAATCCATGGTTCAGCTG
TCTTTGGTTATGAGGGCTGGCTTGCTGGCTACCAGATGACCTTTGACAGTGCCAAATCAAAGCTGACAAGGAATA
ACTTTGCAGTGGGCTACAGGACTGGGGACTTCCAGCTACACACTAATGTCAATGATGGGACAGAATTTGGAGGAT
CAATTTATCAGAAAAGTTTGTGAAGATCTTGACACTTCAGTAAACCTTGCTTGGACATCAGGTACCAACTGCACTC
GTTTTGGCATTGCAGCTAAATATCAGTTGGATCCCACTGCTTCCATTTCTGCTGGTGTGAAGCTTACACTCTCTG
CTCTGGTAGACGGGAAGAGCATTAAATGCTGGAGGCCACAAGGTTGGGCTCGCCCTGGAGTTGGAGGCTTAATCCA
GCTGAAAGAAACCTT

833/6881
FIGURE 776

GAGAAAACGGCCGGGCGGCGGTGGCTGTAGGTTGTGCGGCTGCAGCGGCTCTTCCCTGGGCGGACGATGGACAGC
CAGGGCAGGAAGGTGGTGGTGTGCGACAACGGCACC GGTTTGTGAAGTGTGGATATGCAGGCTCTAACTTTCCA
GAACACATCTTCCCAGCTTTGGTTGGAAGACCTATTATCAGATCAACCACCAAAGTGGGAAACATTGAAATCAAG
GATCTTATGGTTGGTGATGAGGCAAGTGAATTACGATCAATGTTAGAAGTTAACTACCCTATGGAAAATGGCATA
GTACGAAATTGGGATGACATGAAACACCTGTGGGACTACACATTTGGACCAGAGAACTTAATATAGATACCAGA
AATTGTAAATCTTACTCACAGAACCTCCTATGAACCCAAACAAAAACAGAGAGAAGATTGTAGAGGTAATGTTT
GAAACTTACCAGTTTTCCGGTGTATATGTAGCCATCCAGGCAGTTCTGACTTTGTACGCTCAAGGTTTATTGACT
GGTGTAGTGGTAGACTCTGGAGATGGTGTGACTCACATTTGCCAGTATATGAAGGCTTTTCTCTCCCTCATCTT
ACCAGGAGACTGGATATTGCTGGGAGGGATATAACTAGATATCTTATCAAGCTACTTCTGTTGCGAGGATACGCC
TTCAACCACTCTGCTGATTTTGAAACGGTTCGCATGATTAAAGAAAACTGTGTTACGTGGGATATAATATTGAG
CAAGAGCAGAACTGGCCTTAGAAACCACAGTATTAGTTGAATCTTATACACTCCCAGATGGACGTATCATCAA
GTTGGGGGAGAGAGATTGAAGCACCAGAAGCTTTATTTAGCCTCACTTGATCAATGTTGAAGGAGTTGGTGT
GCTGAATTGCTTTTTAACACAATTACGGCAGCTGACATTGATACCAGATCTGAATCTACAAACACATTGTGCTT
TCTGGAGGGTCTACTATGTATCCTGGCCTGCCATCACGGTTGGAACGAGAACTTAAACAGCTTTACTTAGAACGA
GTTTTGAAGGGTGTAGTGGAAAACTTTCTAAATTTAAGATCCGCATTGAAGACCCACCCCGCAGAAAGCACATG
GTATTCCCTGGGTGGTGCAGTTCTAGCGGATATCATGAAAGACAAAGACAACTTTTGATGACCCGACAAGAGTAC
CAAGAAAAGGGTGTCCGTGTGCTAGAGAACTTGGTGTGACTGTTGATAAACTCCAAAGCTTGTCCCGTCATA
CCCCTAATGCTTTCTTTTTCTTTTATTGCCAATCTTTGAACCTCATTCAACTCCAGGACATGGAAGAGGCCTCTC
TCTGCCCTTTGACTGGAAGGTCAAGTTTTATTCTGGTGTCTTGGGGAAGCTTTGTTAAATTTTTGTTAATGTGG
GTAAATCTGAGTTTAATTCAACTGCTTCCCTACATAGACTAGAGGGCTAAGGATTCTGCTGCTGCTTGTCTTCT
TCTAAGTAGGCATTTAGATCATTCTATAGGCTTCTATTTTCACTTTACTGCTCTAATGCTGCTAGTCGTAGTC
TTTAGCACACTAGGTGGTATGCCTTTATTAGCATAAAACAAAAAACTTTAACAGGAGCTTTTACATATTACTG
GGATGGGGGGTGGTTCGGGATGGGTGGGCAGCTGCTGAACCCCTTAGGGCATTTCCTCTGTAATGTGGCGCTTTC
AACTGTACTGCTGCAGCTTTAAGTACCTTAAAGCTTCTCCTGTGAACCTCTTAGGGAAATGTTAGGTTCAGAACT
AAAGTGTTTTGGGTGGGTTTTGTTGCGGGGGGAGGGTAACAATGGGTGGTCTTCTGATTTTTATTTTTGAGGTT
TTGTCAACTGGAGTACGTAGAGGAACTTTATTTACAGTACTTTGATTTGGCAGGTTTTCTTCTACTTGTGCTCTG
CCTGGAGCTGTTTCCATATGATATAAAAAGCAAGTGTAGTATTCCATTACTATGTGGCTTAGGGATTTATTTGTT
TTTTAAATCAACCATGTTAGCTGGGATTAGACTCCCTACAGTCTTCAATGGAAAAGTAACATTTAAAAATCCT
TTGGGTAATTGCAATTACAGATTTAAAAGAGCTTAAGATCTGGTGTGTTTGTAAATGCTTCTGTTTATTCCAGAAG
CATTAAAGTAACCCATTGCCAAGTATCATTCTTGCAAATTATTCTTTATATAACTGACCAGTGCTTAATAAAAC
AAGCAGGTACTTACAAATAATTACTGGCAGTAGGTTATAATTGGTGGTTTAAAAATAACATTGGAATACAGGACT
TGTTGCCAATTGGGTAATTTTCATTAGTTGTTTGTGTTTGTGATTGAAACCTGGAAATACAGTAAATTTGA
CTGTTTAAATGTTGGCCAAAAAATCAAGATTTAATTTTTTTTATTGTACTGAAAACTAATCATAACTGTAA
TTCTCAGCCATCTTTGAAGCTTGAAAGAAGAGTCTTTGGTATTTTGTAAACGTTAGCAGACTTTCCTGCCAGTGT
CAGAAAATCCTATTTATGAATCCTGTCGGTATTCCTTGGTATCTGAAAAAATACCAAATAGTACCATACATGAG
TTATTTCTAAGTTTGAAAAATAAAAAGAAATTGCATCACACTAATTACAAAATA

834/6881
FIGURE 777

ATGGAGACACTTGTACAAGTGAATCTCTTTAGAAAAGAGAGATTTACGCCAGGTACGTGGAACACAGTCTGAGCAG
CACTGCACACCGTATCAAATCAGGTATCCTGTAGGAAGAGTACCTGGGGATGAAATCAACAAGTTCTTAGTGGA
TCTTCTCATTTCTTAGAATGTAATGTGGTTGGATGTTTGGAAATTGGTACGAAAGGGGAGAATCCTTTCCCCCA
TTAGGAAAATGGGGAGTGGGGGCAGAGGAAATGAAGCCCAGGTTCTTCATCGGATGGAAGTTCTCTGGTCAGTGT
ACACAGTTTGCTCTTGAGAGGACTGATGGCCTGGGATATGTTATTGATGGTCCTGAAAGGGATCTGCGTTTGAAT
AGGTCAGAGAAGATGTTCTGGAAGTCTGGAGACTGGGATACGCTCCCTTCCCTACAGGCTCATTATCCAGCGTT
CTGGCAGTACCTCTGGTACCGCACAGACTTCAGGGGACCTTGAACAAGCTCCATTTTGTCTCCTGGAAGCTCAT
TCCACCAGGTTTGAGTATGAGCTTCACCACTCGCTCCACCTTCTCCACCAACTACCACTCCCTGGGCTCTGTCCA
GGTGCCAGCTAAGACGCTCGGCCAGTCAGCAGCATGACCAGTATCTCTGCAGGGGCTGGGGGCTCTGGTTCCCA
GATCTCCATGTCCCCCTCCACCAGCTTCTGGGGTGGCATGGGGTCTGGGGCCCTGGCCGTGGGGATGGCCAGGGT
TCTGGCAGGATGGAAGACATCCAGAACAAGAAGGAGACCATGCAAAGCCTGAATGACTGCCTGGCCTCTACCTG
GACAGAGTGAGGAGACTGGATACCAAGAATCGGAAGCTGGAGAGCAAAATCCGGGAGCACCTGGAGAAGAAGGGA
CCCCAAGTCAGAGACTGGAGCCATTACTTCAAGACCGTGGAGGACCTGAGGGCTCAGATCTTCGCAAATACTGTG
GACAATGCCCCGATCGTTCTGCAAGTCGACAATGCCCATCTTGCTGATGACTTTAGAGTCAAGTATGTGACAGAG
CTGGCCATGCGCCAGTCTATGGAGAGCGACATCCATGGGCTCCGCAAGGTCATTGATGACACCAATGTCACTCGG
CTGCAGCTGGAACAGAGATCGAGGCTCTCAAGGAGGAGCTGCTCTTCATGAAGAAGAACGAAGAAGGGGAAATA
AAAGGCCTACAAGCCCAGATTGCCAGCTCTGGGTTGACTGTGGAGGTAGATATCCCCAAATCTCAGGACCTTGCC
AAGCTCATGGCAGACATCTGGGCCCAATATGATGAGCTGGCTCGGAAGAGCCAAGAGGAGCTGGGCAAGTACTGG
TCTCAGCAGATTGAGGAGAGTATCACAGTAGTCACCATGCAGTCCACCGAGATGGAGCAGTTCAACAGGATCCTG
CTGCACCTGGAGTCAGAGCTGGCACAGACCTGGGCAGAGGGACAGTGCCAGGCCCAGGAGTACGAGGCCCTGCTG
AGCATCAGGGTCAAGCTGGAGGCTGAGATCGCCACCTACCACCACTGCTGGAAGATACCACCACCGGCCGGATA
GTGGATGGCAAAGTGGGGTCTGAGACCAACAACACCAAAGTTCTGAGACATTAA

835/6881

FIGURE 778

METLVQVNLFRKRDLRQVRGTQSEQHCTPYQIRYPVGRVPGDEINKFLVESHFLECNVVGCLEIGTKGENPFPQ
LGKVGVGAEEMKPRFFIGWKFSGQCTQFALERTDGLGYVIDGPERDLRLNRSEKMFWKSGDWDTLPSLQAHSSSV
LAVPSGTAQTSGDLEQAPFCLLEAHSTRFEYELHHSLLHQLPVPGLCPGAQLRRSASQQHDQYLCRGWGLWFP
DLHVPLHQLLGWHGVWGPGRGDGQSGRMEDIQNKKETMQSLNDCLASYLDVRRLDTKNRKLESKIREHLEKKG
PQVRDWSHYFKTVEDLRAQIFANTVDNARIVLQVDNAHLADDFRVKYVTELAMRQSMESDIHGLRKVIDDTNVTR
LQLETEIEALKEELLFMKKNEEGEIKGLQAQIASSGLTVEVDIPKSQDLAKLMADIWAQYDELARKSQEELGKYW
SQQIEESITVVMQSTEMEQFNRIILLHLESELAQTWAEGQCQAQYEALLSIRVKLEAEIATYHHLEDTTTGRI
VDGKVGSETNNIKVLRH

836/6881
FIGURE 779A

CGGTCCCTCGCCGCGCCGCCCGAGGGGCACTTCCGGCGGCGGTTCACTTCCTGGTTGGGTGGATGGAGCCGGGG
GGGAGCGCGCGCGGGGGAGGGGCGCGGGTCAGTCTCGGCCCGGCGCTCCCGGGATCAGCTGGCGGGCGGGCGGG
AGCCGAGCGCGCGCCCGGCTCTCGCTGCAGCGCCGCTCTTCTCTCGCTCGCAGGCCGCGCCGCGGGCGCGGTGACA
ATGTCGCGGGGCTGGTAGCAGGGCGCGGCCGCGGAGCCGCTCTCAAGTTTAACTTACACGAATCGCTTTCTGGA
GGAGGAGGGGACCGCTGCGCGATTGACACGCATATTCCTATAGGCATCCTCCCTCAGCCCCACCCCCACGGCC
GGATTGCGGTGGCTCCTCTCCGAGGTGAAATCTGAGAAGAAATCCTTGATCTCTTTTCTTAAAAAAAAAAAAA
AAAAAAAAAATCTAGAAACCATCGGTATTTTGCTTTGCTGCTCCCTATTTCGCAAGATGAAGAAGTTTTTCGACTC
CCGGCGAGAGCAGGGCGGCTCTGGCTGGGCTCCGGCTCCAGCGGAGGAGGGGCGAGCACCTCGGGCCTGGGCAG
TGGCTACATCGGAAGAGTCTTCGGCATCGGGCGACAGCAGGTACAGTGGACGAGGTGTTGGCGGAAGGTGGATT
TGCTATTGTATTTCTGGTGAGGACAAGCAATGGGATGAAATGTGCCTTGAAACGCATGTTTGTCAACAATGAGCA
TGATCTCCAGGTGTGCAAGAGAGAAATCCAGATAATGAGGGATCTTTCAGGGCACAAGAATATTGTGGGTACAT
TGATTCTAGTATCAACAACGTGAGTAGCGGTGATGTATGGGAAGTGCTCATTCTGATGGACTTTTGTAGAGGTGG
CCAGGTGGTAAACCTGATGAACAGCGCCTGCAACAGGCTTTACAGAGAATGAAGTGCTCCAGATATTTTGTGA
TACCTGTGAAGCTGTGCGCGCTGCATCAGTGCAAACTCCTATTATCCACCGGGACCTGAAGGTTGAAAACAT
CCTCTTGATGACCGAGGCCACTATGTCCTGTGTGACTTTGGAAGCGCCACCAACAAATTCAGAATCCACAAC
TGAGGGAGTCAATGCAGTAGAAGATGAGATTAAGAAATACACAACGCTGTCTATCGAGCACCAGAAATGGTCAA
CCTGTACAGTGGCAAAATCATCACTACGAAGGCAGACATTTGGGCTCTTGATGTTTGTGTATAAATTATGCTA
CTTCACTTTGCCATTTGGGGAAAGTCAGGTGGCAATTTGTGATGGAACTTCACAATTCCTGATAATTCTCGATA
TTCTCAAGACATGCACTGCCTAATTAGGTATATGTTGGAACAGACCCTGACAAAAGGCCGGATATTTACCAGGT
GTCTACTTCTCATTTAAGCTACTCAAGAAAGAGTGCCCAATTCCAAATGTACAGAACTCTCCCATTCCTGCAAA
GCTTCTGAACCACTGAAAGCCAGTGAGGCAGCTGCAAAAAGACCCAGCCAAAGGCCAGACTGACAGATCCCAT
TCCCACCACAGAGACTTCAATTGCACCCCGCCAGAGGCCATAAGCTGGGCAGACTCAGCCGAACCCAGGAATCCT
TCCCATCCAGCCAGCGCTGACACCCCGGAAGAGGGCCACTGTTAGCCCCCACCCTCAGGCTGCAGGATCCAGCAA
TCAGCCTGGCCTTTTAGCCAGTGTTCCCAACCAAAACCCCAAGCCCCACCCAGCCAGCCTCTGCCGCAAACTCA
GGCCAAGCAGCCACAGGCTCCTCCCACTCCACAGCAGACGCTTCTACTCAGGCCCAGGGTCTGCCGCTCAGGC
CCAGGCCACACCCACAGCACCAGCAGCAACTCTTCTCAAGCAGCAACAGCAGCAGCAACAGCCACCGCCAGCACA
GCAGCAGCCGGCAGGCACGTTTTTACCAGCAGCAGCAGGCCCAGACTCAGCAGTTTTCAGGCAGTACATCCAGCAAC
CCAGAAACAGCAATTGCTCAGTTCCCTGTGGTGTCCTCAAGGAGGCTCTCAACAGCAGCTAATGCAGAATTTCTA
CCAGCAGCAGCAGCAGCAGCAACAACAACAGCAACAGCTGGCCACAGCCCTGCATCAACAACAGCTGAT
GACTCAGCAGGCTGCCTTGAGCAAAAGCCCACTATGGCAGCAGGACAGCAGCCCCAGCCACAGCCAGCTGCAGC
CCCACAGCCAGCCCCTGCCAGGAGCCAGCGATTCAAGCCCCAGTAAGACAACAGCCAAAGGTTGAGACAACCCC
ACCTCCTGCCGTCCAGGGGCGAGAAAGTTGGATCTCTCACTCCACCCTCATCCCCAAAACCAACGTGCTGGGCA
CAGGCGTATTCTCAGTGACGTAACCCACAGTGCACTTTGGGGTCCCTGCCAGCAATCAACCCAGCTGCTCCA
GGCAGCTGCAGCTGAGGCCAGTCTCAATAAGTCCAAGTCTGCAACCACCACTCCATCAGGCTCTCCTCGGACCTC
TCAACAAAACGTTTATAATCCTTCAGAAGGCTCTACGTGGAATCCCTTTGATGACGATAATTTCTCAAACCTCAC
AGCTGAAGAACTGCTAAACAAGGACTTTGCCAAGCTTGGGAAGGCAACATCCCGAGAAGCTTGGAGGCTCAGC
TGAGAGTTTGATCCAGGCTTTCAATCAACCCAAGGTGATGCTTTTGCTACGACCTCATTTTCTGCTGGAACCTGC
TGAAAAAAGGAAGGTGGGCAGACTGTGGACTCTGGCCTCCCGCTTCTAAGCGTGTCTGATCCTTTCATTCTCT
TCAAGTACCTGATGCACCAGAAAACTAATTGAGGGACTCAATCTCCTGACACTTCTCTTCTGCTCCCTGACCT
CTTGCTATGACAGATCCTTTTGGTAGCACTTCTGATGCTGTAATTGGTAAAGTCATCATCTCTGTTTCTTCAGT
CATGCATGATATGTGTGCTGTTTCAAGAATGACAAGTACCTAGTTAACCAATCCCTGGGGAATAGCCCTGCCAC
CCCAGAAGCCAAGGCTATTTAATCTCACTTGATCACTCCAAATGAAGTGTTTTCTTGCTTTTCGGGGGTACAA
CGTCTATTTTTTGCTTTCTTGATATCTGGAGATTTCTAGAGTGGATCTCTTATGAATGAGGAGGAATGTGAAA
GTTTCACATTTAGTTAGAAGAACTTTAAAAAATGGTTTCTAACTAGACTACCTTTACTGAACCTTAATGAAATTT
AGCAGATTCTTCTTTTTTTTTTTTTTGTGTTGAGACAGAGTTTCACTCTTGTGCTAGGCTGGAGTGAATGGCA
AGATCTCAGCTCACTGCAACCTCCATCTCCCGGTTCAAGCAATCTCCTGCCTCAGCCTCCCGAGTAGCTGAGA
TTACAGGCATGCACCACCATGCCTGGCTAATTTTATATTTTAGTAGAGATAGGGTTTCTCCATGTTGGTCAGG
CTGGTCTCGAACTCCTGACCTCAGGTGATCCACCTGCCTTGGCCTCCCAAGTGCTGGGATTACAGGGGTGAGCC

837/6881
FIGURE 779B

ACCATGCCCAGCCCAGATTCTTCTTTAGAAAGGCCCATGTACATCCAGCTGCAGTGGGACGAGCCTGGGAAGCATA
GGAAGACCCCATCCCTACAAAAAAATGTTGTTAGTTAGCCAGGTGTGGGGGTGCACACCTGTGCTACCAGCTCT
TTGGGAGGCTGAGACGGAAGATCACTTTGAGCCAGGGGGTTTGAGGCTGCAGTGAGCCCTGATCATGCCACTGCC
CTCCAGCTTAAGTGACAGAGACCCATCTTAAACAAAATTAAAAAATTTAAAAAGGTCCATGTATATACCAAGTAT
CAGTAACTGAGAAAACATTTTGCCCACTGGTTAAGCCCCATAAATAGAATCCAGGGCAGGGGCTTGGTCTGCTTC
CTCTCTCACATTTTCCTATGTTCCAACAGCAAGGCAGGCAAGATTTCAGAAAATGTCCAAGGCCCTGTACTCCAG
GAGGGTCCAATCATGAGGCACCTAATAATGCCTTTGGGCAGAAAGTCCTAGAACTCAGAAGTGCCATGACCTTGC
ATAGGGGTAGGCCTGCAGCTGAGGGGTGTCACGTGGTGCAAAAGGCCTCCAGCTACATCCTGGGATGGGGTTTG
AGATGGATTAATGGCCTTTGTCATGGCACAGAGGATGTTTGCCCCCAGGTTTATCCCTTAATAAAAGTGAAATTAC
CAGAAAAAATTAAGGTGCCTTTCAGCCCATCTCTTCTGACATGGAAGAGGTTTGCTGAGCGGCTTATGTTCTGAA
TTTGTATTATGAGAACTGATCATTAGTGAGACTGGCCACAGTATTTAACCTTGACATGCATGCTCAGTGAAGCCT
AGACATTGAGAGCAGCAGAAATAAAGTAACTTATTTTCTTTCCACCTTGCTGAGACTGGCATTTTAGAGACCT
GTTAAAAGAAAGTCTTCACATGGCTGGGTGCAGTGGCTCACACTTGTAACACAGGACTTTGGGAGGCCAAGGTG
GGATAGATCGCTTGAGCTCAGGAGCAAGACACCATCTCAAAAAATGAGAATT

838/6881
FIGURE 780

GCCCCAGGTGCGCTTCCCCTAGAGAGGGATTTTCCGGTCTCGTGGGCAGAGGAACAACCAGGAACCTTGGGCTCAG
TCTCCACCCCAACAGTGGGGCGGATCCGTCCCGGATAAGACCCGCTGTCTGGCCCTGAGTAGGGTGTGACCTCCGC
AGCCGCAGAGGAGGAGCGCAGCCCGGCTCGAAGAAGTCTGTCTGGGTGGCTGAACTCTGATCTTGACCTAGAG
TCATGGCCATGGCAACCAAAGGAGGTACTGTCAAAGCTGCTTCAGGATTCAATGCCATGGAAGATGCCCAGACCC
TGAGGAAGGCCATGAAAGGGCTCGGCACCGATGAAGACGCCATTATTAGCGTCCTTGCCCTACCGCAACACCGCCC
AGCGCCAGGAGATCAGGACAGCCTACAAGAGCACCATCGGCAGGGACTTGATAGACGACCTGAAGTCAGAAGTGA
GTGGCAACTTCGAGCAGGTGATTGTGGGGATGATGACGCCCACGGTGTCTGTATGACGTGCAAGAGCTGCGAAGGG
CCATGAAGGGAGCCGGCACTGATGAGGGCTGCCTAATTGAGATCCTGGCCTCCCGGACCCCTGAGGAGATCCGGC
GCATAAGCCAAACCTACCAGCAGCAATATGGACGGAGCCTTGAAGATGACATTGCTCTGACACATCGTTTCATGT
TCCAGCGAGTGTGGTGTCTCTGTGACGTGGTGGGAGGGATGAAGGAAATTATCTGGACGATGCTCTCGTGAGAC
AGGATGCCCAGGACCTGTATGAGGCTGGAGAGAAGAAATGGGGGACAGATGAGGTGAAATTTCTAACTGTTCTCT
GTTCCCGGAACCGAAATCACCTGTTGCATGTGTTTGATGAATACAAAAGGATATCACAGAAGGATATTGAACAGA
GTATTAAATCTGAAACATCTGGTAGCTTTGAAGATGCTCTGCTGGCTATAGTAAAGTGCATGAGGAACAAATCTG
CATATTTTGTCTGAAAAGCTCTATAAATCGATGAAGGGCTTGGGCACCGATGATAACACCCTCATCAGAGTGATGG
TTTCTCGAGCAGAAATTGACATGTTGGATATCCGGGCACACTTCAAGAGACTCTATGGAAAGTCTCTGTACTCGT
TCATCAAGGGTGACACATCTGGAGACTACAGGAAAGTACTGCTTGTTCTCTGTGGAGGAGATGATTAAAATAAAA
ATCCCAGAAGGACAGGAGGATTCTCAACACTTTGAATTTTTTAACTTCATTTTCTACACTGCTATTATCATT
TCTCAGAATGCTTATTTCCAATTAACGCCTACAGCTGCCTCCTAGAATATAGACTGTCTGTATTATTATCAC
CTATAATTAGTCATTATGATGCTTTAAAGCTGTACTTGCAATTTCAAAGCTTATAAGATATAAATGGAGATTTTAA
AGTAGAAATAAATATGTATTCCATGTTTTTAAAGATTACTTTCTACTTTTGTGTTTCACAGACATTGAATATATT
AAATTATTCCATATTTTCTTTTTCAGTGAAAAATTTTTTAAATGGAAGACTGTTCTAAAATCACTTTTTTCCCTAA
TCCAATTTTATAGAGTGGCTAGTAGTTTCTTCATTTGAAATTTGAAGCATCCGGTCAGTAAGAATGCCCATCCAGT
TTTCTATATTTATAGTCAAAGCCTTGAAAGCATCTACAAATCTCTTTTTTTAGGTTTTGTCCATAGCATCAGTT
GATCCTTACTAAGTTTTTTCATGGGAGACTTCCTTCATCACATCTTATGTTGAAATCACTTTCTGTAGTCAAAGTA
TACCAAACCAATTTATCTGAACTAAATTTCTAAAGTATGGTTATACAAACCATATACATCTGGTTACCAAACATA
AATGCTGAACATTCCATATTATTATAGTTAATGTCTTAATCCAGCTTGCAAGTGAATGGAAAAAATAAGCTT
CAAAGTAGGTATTCTGGGAATGATGTAATGCTCTGAATTTAGTATGATATAAAGAAAACTTTTTGTGCTAAAAA
TACTTTTTAAATCAATTTTGTGATTGTAGTAATTTCTATTGCACTGTGCCTTTCAACTCCAGAAACATTCTG
AAGATGTACTTGGATTTAATTAAGTTCACTTTGT

839/6881
FIGURE 781

MAMATKGGIVKAASGFNAMEDAQTLRKAMKGLGTDEDAIISVLAYRNTAQRQEIRTAYKSTIGRDLIDDLKSELS
GNFEQVIVGMMTPTVLYDVQELRRAMKGAGTDEGCLIEILASRTPEEIRRISQTYQQYGRSLEDDIRSDTSFMF
QRVLVSLSAGGRDEGNYLDDALVRQDAQDLYEAGEKKWGTDEVKFLTVLCSRNRNHLHVFDEYKRISQKDIEQS
IKSETSGSFEDALLAIVKCMRNKSAYFAEKLYKSMKGLGTDDNTLIRVMVSRAEIDMLDIRAHFKRLYGKSLYSF
IKGDTSGDYRKVLLVLCGGDD

840/6881
FIGURE 782

CGCCACCGCCGCGCCCTCGCCACCCGCCCGCCCGCCGCTCCCGGCCCGCTCGCCCCCTCCGCGCCGCGCCCGCC
GCCCCCTGCGACTACGCTGCGGCCCTCCCGCCCGCTCCCGCTCGCTCCCGCGGCCCTCGCTCGCCTCGCGCCGGCAG
TTTTGGGCCTACACCTCCCCTCCCCCGCCAGCCGCCAAAGACTTGACCACGTAACGAGCCCAACTCCCCGAAC
GCCGCCCCGCGCTCGCCATGGATGCCGGTGTGACTGAAAGTGGACTAAATGTGACTCTCACCATTGCGCTTCTTA
TGCACGGAAAGGAAGTAGGAAGCATCATTGGGAAGAAAGGGGAGTCGGTTAAGAGGATCCGCGAGGAGAGTGGCG
CGCGGATCAACATCTCGGAGGGGAATTGTCCGGAGAGAATCATCACTCTGACCGGCCCCACCAATGCCATCTTTA
AGGCTTTCGCTATGATCATCGACAAGCTGGAGGAAGATATCAACAGCTCCATGACCAACAGTACCGCGGCCAGCA
GGCCCCCGGTACCCCTGAGGCTGGTGGTGCCGGCCACCCAGTGCGGCTCCCTGATTGGGAAAGGCGGGTGTAAAG
TCAAAGAGATCCGCGAGAGTACGGGGGCGCAGGTCCAGGTGGCGGGGGATATGCTGCCCAACTCCACCGAGCGGG
CCATCACCATCGCTGGCGTGCCGCGAGTCTGTCAACGAGTGTGTCAAGCAGATTTGCCTGGTCATGCTGGAGACGC
TCTCCCAGTCTCCGCAAGGGAGAGTCATGACCATTCCGTACCAGCCCATGCCGGCCAGCTCCCCAGTCATCTGCG
CGGGCGGCCAAGATCGGTGCAGCGACGCTGCGGGCTACCCCCATGCCACCCATGACCTGGAGGGACCACCTCTAG
ATGCCTACTCGATTCAAGGACAACACACCATTTCTCCGCTCGATCTGGCCAAGCTGAACCAGGTGGCAAGACAAC
AGTCTCACTTTGCCATGATGCACGGCGGGACCGGATTGCGCCGAATTGACTCCAGCTCTCCAGAGGTGAAAGGCT
ATTGGGCAAGTTTGATGTCATCTACTCAAACCACCCATGAACTCACCATTCCAAATAACTTAATTGGCTGCATAA
TCGGGCGCCAAGGCGCCAACATTAATGAGATCCGCCAGATGTCCGGGGCCAGATCAAAATTGCCAACCCAGTGG
AAGGCTCCTCTGGTAGGCAGGTTACTATCACTGGCTCTGCTGCCAGTATTAGTCTGGCCCAGTATCTAATCAATG
CCAGGCTTTCCTCTGAGAAGGGCATGGGGTGCAGCTAGAACAGTGTAGGTTCCCTCAATAACCCCTTCTGCTGT
TCTCCCATGATCCAACCTGTGTAATTTCTGGTCAGTGATTCCAGGTTTTAAATAATTTGTAAGTGTTCAGTTTCTA
CACAACTTTATCATCCGCTAAGAATTTAAAAATCACATTCTCTGTTTACGCTGTTAATGCTGGGATCCATATTTAG
TTTTATAAGCTTTTCCCTGTTTTTGTGTTTTGGGTTTTTGGCTCATGAATTTTATTTCTGTTTGTGCGATA
AGAAATGTAAGAGTGGAAATGTTAATAAATTTAGTTTCTGTAATGTCAAGAATTTAAGAATTAATAAACGG
ATTGGTTAAAAAATGCTTCATATTTGAAAAAGCTGGGAATTGCTGTCTT

841/6881
FIGURE 783

CCAAGCTGAACCAGGTGGCAAGACAACAGTCTCACTTTGCCATGATGCACGGCGGGACCGGATTGCCCGGAATTG
ACTCCAGCTCTCCAGAGGTGAAAGGCTATTGGGCAAGTTTGGATGCATCTACTCAAACCACCCATGAACTCACCA
TTCCAAATAACTTAATTGGCTGCATAATCGGGCGCCAAGGCGCCAACATTAATGAGATCCGCCAGATGTCCGGGG
CCCAGATCAAATTGCCAACCCAGTGGGAAGGCTCCTCTGGTAGGCAGGTTACTATCACTGGCTCTGCTGCCAGTA
TTAGTCTGGCCCAGTATCTAATCAATGCCAGGCTTTCCTCTGAGAAGGGCATGGGGTGCAGCTAGAACAGTGTAG
GTTCCCTCAATAACCCCTTTCTGCTGTTCTCCCATGATCCAACGTGTGTAATTTCTGGTCAGTGATTCCAGGTTTT
AAATAATTTGTAAGTGTTTCAGTTTCTACACAACCTTTATCATCCGCTAAGAATTTAAGCTTTTCCTGTTTTAGT
TTTGTTTTGGGTTTTTTGGCTCATGAATTTTA

842/6881
FIGURE 784

CGCAGAGCTCGCGCGTCGGTGTGGTTACAGCCGGCAGCCACGGCACCTCCTTCCGGCCGACTAGTCTCCAGGTCC
CGCGGTCTGGCCCCCTGGTCACCCGGCGCTGTCCCTCGTTCCTTGAAGATCTCCAGTGCTACCTTTTGTGTAACAT
CTCGAATACACCAAACCTTCTAGCTGCTAATACTCAGATTTATGGACAAGAAGTTATCATTGAAATTAAATGGTG
GCAGACATGTCCAAGGAATATTGCGGGGATTTGATCCCTTATGAACCTTGTGATAGATGAATGTGTGGAGATGG
CGACTAGTGGACAACAGAACAATATTGGAATGGTGGTAATACGAGGAAATAGTATCATCATGTTAGAAGCCTTGG
AACGAGTATAAATAATGGCTGTTCAGCAGAGAAACCCATGTCCTCTCTCCATAGGGCCTGTTTTACTATGATGTA
AAAATTAGGTCATGTACATTTTTCATATTAGACTTTTTGTAAATAAACTTTTGTAATAGTCAAAAATGCTTTCTC
AGATGTTCTGAATATAGAATATCAGCTCTCATTCCAGTTTTTTCTAACATGAATTTTCCTGGTTGACATTGATTT
CAAAGGGTTTTATGCATTAAAGTGAAAGAATCTTATTAAATGTGAAACATGGCAAGGATTCCTTCTGTCCATCAT
CTTCAACACTGAACAGATAAAGAAGTTACAAACATGAGTGAGTTTCTTCTAAAACACTAATTTTC

843/6881
FIGURE 785

GTACGCCGATTCCATATGGGCGCCGGCGCGGAGCGCCGCGGGGCGAGCGCGGGGTCGCCATGGCTGAGCTGCAGCA
GCTCCGGGTGCAGGAGGCGATGGAGTCCATGGTGAAGAGTCTGGAAAGAGAGAACATCCGGAAGATGCAGGGTCT
CATGTTCCGGTGCAGCGCCAGCTGTTGTGAGGACAGCCAGGCCTCCATGAAGCAGGTGCACCAAGTGCATCGAGCG
CTGCCATGTGCTCTGGCTCAAGCCCAGGCTTTGGTCAACAGTGAAGCTGGAGAAGTTCAGGACCGCCTGGCCCCG
GTGCACCATGCAATGCAATGACAAAGCCAAAGATTCAATAGATGCTGGGAGTAAGGAGCTTCAGGTGAAGCAGCA
GCTGGACAGTTGTGTGACCAAGTGTGTGGATGACCACATGCACCTCATCCCACTATGACCAAGAAGATGAAGGA
GGCTCTCTTATCAATTGAAAAATAAAGTATTTGCCAGTGGCCATCAGGCTGAGGGCAAGAATATATTTTTTATA
AGGAATTGGGAATTTTAGTCTTTTAAGCAAAGTTTACGAATGAAGAAATGAAGGATGGCCACAAGCGTAAGGCAT
AIGTCACTTGCTCTGGACACTGGTTATTTTATGTTTCAGTCCCTAAAAAATGAAATGGAAAAAAGTGGTGCTAA
ATCGAGTCAGAGATATTACAGGAGAGTTTTAGAGCTTATTATTTCTGTGGCCAGTGTCTGTCTGGCAGTAAGG
CTCTCCCTGTAAACAAGCCAGAGCCCTCCAAGGTACCAGACTCTTCTTACTACACAGGTACTAACAGGCTGGCAG
GTTAGAGTTGGTGGAGTCTGAGGAGAGATATTTTCTTTGTTGCCAACATCCTGTTTACCAAAAGTGTCAACCC
ACCATCTTCCATAAGCTGTGAAACAAAATCAATGAGGTCACTAACTTAGAAGGGAAGAAAGTTTTCTGGGTCTT
TGTTTTCTTGATTGGGGTAATTTATACAAGGGCATAACAAGTTGATTTAAGATGTGGAAGTGGGAGGTAGACTA
GTTTGGATAAGAACTTTGAAATGTTCTTGTGGATCCCCATTTCTGGTCATCAAGATGTGGATGTACATTTCTTA
AAATTATTACATGCTGCATCTTTCAGCCTGGAGACTGTGCAGAAACATGAGAGGTGATGACACACTAATTATGGG
AAGCAGAATTACTGGCTGATGGCCCTGAGGCTGTGTGTAACAAAATGACAGGACAATCTTGACAGTAACACTTTC
CCCTTGAAGAGAAGGGGGTTTTGATTGTGATATATACTAGTATCTAGGAATGAACAGTAAAGAGGAGCAGTTGG
CTACTTGATTACAACAGAGTAAATGAAGTACTGGATTGGGAAAACCTGGTTTTATTAGAACATATGGAATGAAA
GCCTACACCTAGCATTGCCTACTTAGCCCCCTGAATTAACAGAGCCCAATTGAGACAAACCCCTGGCAACAGGAA
ATTCAAGGGAGAAAAAGTAAGCAACTTGGGCTAGGATGAGCTGACTCCCTTAGAGCAAAGGAGAGACAGCCCCCA
TTACCAAATACCATTTTTGCCTGGGGCTTGTGCAGCTGGCAGTGTTCCTGCCCCAGCATGGCACCTTATTGTTTT
GATAGCAACTTCGTTGAATTTTACCAACTTATTACTTGAAATTATAATATAGCCTGTCCGTTTGCTGTTTCCAG
GCTGTGATATATTTTCTAGTGGTTTGACTTTAAAAATAAATAAGGTTTAATTTTCTCCCC

844/6881
FIGURE 786

MAELQQLRVQEAMESMVKSLEARENIRKMQGLMFRCSASCCEDSQASMKVHQCIERCHVPLAQAQALVTSELEKF
QDRLARCTMHCNDKAKDSIDAGSKELQVKQQLDSCVTKCVDDHMHLIPTMTKKMKEALLSIGK

845/6881
FIGURE 787A

CAGACCCGCCCGTGGCTGCAGCAGTGGTGTCCCATTTTAAATGACTGCCAGATTCCCACACTCAGTTCTGCTTCC
ATGGAACCTGCAGGTTTTTGGTGCAGGAGGACAAGCCAGCATGTGTCTGCCATTCTGGGTACGTTGGTGCACGCT
GTGAGCATGCGGACCTCCTGGCCGTGGTGGCTGCCAGCCAGAAGAAGCAGGCCATCACCGCCTTGGTGGTGGTCT
CCATCGTGGCCCTGGCTGTCTTATCATCACATGTGTGCTGATACACTGCTGCCAGGTCCGAAAACACTGTGAGT
GGTGGCGGGCCCTCATCTGCCGGCACGAGAAGCCAGCGCCCTCCTGAAGGGAAGAACCCTTGGTGGCACTCAG
AAACAGTGGTCTGAAGAGCCAGAGGAGGAGTTTGGCCAGGTGGACTGTGGCAGATCAATAAAGAAAGGCTTCTT
CAGGACAGCACTGCCAGAGATGCCTGGGTGTGCCACAGACCTTCCTACTTGGCCTGTAATCACCTGTGCAGCCTT
TTGTGGGCTTCAAACTCTGTCAAGAACTCCGTCTGCTTGGGGTTATTAGTGTGACCTAGAGAAGAAATCAGC
GGACCAGATTTCAAGACTTGTAAAAAAGAACTGCAAGAGACGGACTCCTGTTACCTAGGTGAGGTGTGTGC
AGCAGTTGGTGTCTGAGTCCACATGTGTGCAGTTGTCTTCTGCCAGCCATGGATTCCAGGCTATATATTTCTTTT
TAATGGGCCACCTCCCCACAACAGAATTCTGCCAACACAGGAGATTTCTATAGTTATTGTTTTCTGTCATTTGC
CTACTGGGAAGAAAGTGAAGAGGGGAAACTGTTAATATCACATGAAGACCCTAGCTTTAAGAGAAGCTGTAT
CCTCTAACACGAGACCCCTCAACCAGCCCAACATCTTCCATGGACACATGACATTGAAGACCATCCCAAGCTATC
GCCACCCTTGGAGATGATGTCTTATTTATTAGATGGATAATGGTTTTATTTTTAATCTCTTAAGTCAATGTAAAA
AGTATAAAACCCCTTCAGACTTCTACATTAATGATGTATGTGTGCTGACTGAAAAGCTATACTGATTAGAAATG
TCTGGCCTCTTCAAGACAGCTAAGGCTTGGGAAAAGTCTTCCAGGGTGCGGAGATGGAACCAGAGGCTGGGTAC
TGGTAGGAATAAAGGTAGGGGTTTCAAGAAATGGTGCCATTGAAGCCACAAAGCCGGTAAATGCCTCAATACGTTCT
GGGAGAAAACCTTAGCAAATCCATCAGCAGGGATCTGTCCCTCTGTTGGGGAGAGAGGAAGAGTGTGTGTCTA
CACAGGATAAAACCAATACATATTGTACTGCTCAGTGATTAAATGGGTTCACTTCTCGTGAGCCCTCGGTAAGT
ATGTTTAGAAATAGAACATTAGCCACGAGCCATAGGCATTTTCAAGCCAAATCCATGAAAGGGGACAGTCATTT
ATTTTCCATTTTGTGCTTGGTTGGTTTGTGTGCTTATTTTTTAAAGGAGAAGTTTAACTTTGCTATTTATTTT
GAGCACTAGGAAAATATTCCAGTAATTTTTTTTTCTCATTTCATTGAGGATGCCGGCTTTATTAACAAAAAC
TCTAACAAGTCACCTCCACTATGTGGGTCTTCCCTTCCCTCAAGAGAAGGAGCAATTGTTCCCTTGAGCATCTG
GGTCCATCTGACCATGGGGCCTGCCTGTGAGAAACAGTGGGTCCCTTCAAATACATAGTGATAGCTCATCCCT
AGGAATTTTCATTAATAATTTGGAAACAGAGTAATGAAGAAATAATATATAAACTCCTTATGTGAGGAAATGCTAC
TAATATCTGAAAAGTGAAGATTTCTATGTATTAACCTTAAAGTGACCTAGCTTATTACATCGTGAAAGGTACA
TTTAAATATGTTAAATTTGGCTTGAAATTTTCAAGAGAATTTGTCTTCCCTAATTCTTCTTCTTGGTCTGGAA
GAACAATTTCTATGAATTTTCTCTTTATTTTTTTTATAATTCAAGACAATCTATGACCCGTGTCTTCTTCTT
GCACTCTTATTTAACAATGCCACACCTGAAGCACTTGGATCTGTTTCAAGAGCTGACCCCTAGCAACGTAGTTGAC
ACAGCTCCAGGTTTTTAAATTAATAAGTTCAAGTTTACATCCCTTGGGCCAGATATGTGGGTTGAGGCTT
GACTGTAGCATCTGCTTAGAGACCAATCAACGGACACTGGTTTTTAGACCTCTATCAATCAGTAGTTAGCATCC
AAGAGACTTTGACAGAGCGTAGGAATGAGGCTGGACAGATGGCGGAAGCAGAGGTTCCCTGCGAAGACTTGAGAT
TTAGTGTCTGTGAATGTTCTAGTTCCTAGGTCCAGCAAGTCACACCTGCCAGTGCCTCATCCTTATGCCTGTAA
CACACATGCAGTGAGAGGCCTCACATATACGCCTCCCTAGAAGTGCCTTCCAAGTCAGTCTCTTGGAAACCAGCA
GGTCTGAAAAGAGGCTGCATCAATGCAAGCCTGGTTGGACCATGTCCATGCCTCAGGATAGAACAGCCTGGCT
TATTTGGGGATTTTTCTTCTAGAAATCAAATGACTGATAAGCATTTGGATCCCTCTGCCATTTAATGGCAATGGTA
GTCTTTGGTTAGCTGCAAAAATACTCCATTCAAGTTAAAAATGCATCTTCTAATCCATCTCTGCAAGCTCCCTG
TGTTTCCCTTGCCCTTTAGAAAATGAATTTTCACTACAATTAGAGAATCATTTAACATCCTGACCTGGTAAGCTG
CCACACACCTGGCAGTGGGGAGCATCGCTGTTTCCAATGGCTCAGGAGACAATGAAAAGCCCCCATTTAAAAAAA
TAACAAACATTTTTTAAAGGCCTCCAATACTCTTATGGAGCCTGGATTTTTTCCCACTGCTCTACAGGCTGTGAC
TTTTTTAAGCATCTGACAGGAAATGTTTTCTTCTACATGGAAGATAGACAGCAGCCAACCTGATCTGGAAG
ACAGGGCCCCGGCTGGACACACGTGGAACCAAGCCAGGATGGGCTGGCCATTGTGTCCCCGACAGGAGATGGG
CAGAAATGGCCCTAGAGTTCTTTTCCCTGAGAAAGGAGAAAAAGATGGGATTGCCACTACCCACCCACACTGGTA
AGGGAGGAGAATTTGTGCTTCTGGAGCTTCTCAAGGGATTGTGTTTTGCAGGTACAGAAAACCTGCTGTATCTT
CAAGCCAGGTTTTTCGAGGGCACATGGGTACCAGTTGCTTTTTTCAGTCAATTTGGCCGGGATGGACTAATGAGGC
TCTAACACTGCTCAGGAGACCCCTGCCCTCTAGTTGGTTCTGGGCTTTGATCTTCCAACCTGCCAGTCACAG
AAGGAGGAATGACTCAAATGCCCCAAAACCAAGAACACATTGCAGAAGTAAGACAAAACATGTATATTTTTAAATGT
TCTAACATAAGACCTGTTCTCTCTAGCCATTGATTTACCAGGCTTTCTGAAAGATCTAGTGGTTCACACAGAGAG

846/6881

FIGURE 787B

AGAGAGAGTACTGAAAAAGCAACTCCTCTTCTTAGTCTTAATAATTTACTAAAATGGTCAACTTTTCATTATCTT
TATTATAATAAACCTGATGCTTTTTTTTAGAACTCCTTACTCTGATGTCTGTATATGTTGCACTGAAAAGGTTAA
TATTTAATGTTTTAATTTATTTTGTGTGGTAAGTTAATTTTGATTTCTGTAATGTGTTAATGTGATTAGCAGTTA
TTTTCCTTAATATCTGAATTATACTTAAAGAGTAGTGAGCAATATAAGACGCAATTGTGTTTTTCAGTAATGTGC
ATTGTTATTGAGTTGTACTGTACCTTATTTGGAAGGATGAAGGAATGAATCTTTTTTTCCT

847/6881

FIGURE 788A

[illegible]

848/6881
FIGURE 788B

TCCCCTGACCTTCCCTCACCCCCAGAAACACTTGCTTACAGACCGAAACTGGCATCTTACTCTTGGCACCTTGAC
TTGCACCCTCTGAGGTTCCAACCTCAGTCATTCTTTGTCCAGCAGAGGAGAATCAGAAATGAGCCCTTCAGGATTA
ATCCTCTTGCACCAGCTCTCAGAGAAATGCTGGGTATCCCTGTCCTTGTCCTATCTGTCCATCCTGGGGCCTGG
TAATGGCCACAGTTATTGTTTTAAATGCCAACACTGTCTTCTCATGTTCTTCCGTGGGGCATTGATTAATGAGCA
TTGTTGGCTCCTAAAAATTAGACAATCCATTCTCTTG

849/6881
FIGURE 789

GGCACGAGGCGGAGAACAGGAGCATGGAGGGCGGGCTGGGGCGTGCTGTGTGCTTGCTGACCGGGGCCTCCCGCG
GCTTCGGCCGGACGCTGGCCCCGCTCCTGGCCTCGCTGCTGTCGCCCGGCTCCGTGCTTGTCCTTAGCGCCCGCA
ACGACGAGGCACTGCGCCAGCTGGAGGCCGAGCTGGGCGCCGAGCGGTCTGGCCTGCGCGTGCTGGGGTGCCCG
CCGACCTGGGCGCCGAGGCCGGCTTGACAGCAGCTGCTCGGCGCCCTGCGCGAGCTCCCCCGGGCCCAAGGGGCTGC
AGCGACTGCTGCTTATCAACAACGCGGGCTCTCTTGGGGATGTGTCCAAAGGCTTCGTGGACCTGAGTGACTCCA
CTCAAGTGAACAACACTGAGGCACTGAACTTGACCTCCATGCTCTGCCTGACTTCCAGCGTCCTGAAGGCCTTCC
CGGACAGTCCTGGCCTCAACAGAACCGTGTTAAACATCTCGTCCCTCTGTGCCCTGCAACCTTTCAAAGGCTGGG
CGCTGTACTGTGTCAGGAAAGGCTGCTCGTGATATGCTGTTCCAGGTCCTGGCGCTGGAGGAACCTAATGTGAGGG
TGCTGAACTATGCCCCAGGTCCTCTGGACACAGACATGCAGCAGTTGGCCCGGGAGACCTCCGTGGACCCAGACA
TGCGAAAAGGGCTGCAGGAGCTGAAGGCAAAGGGGAAGCTGGTGGATTGCAAGGTGTCAGCCCAGAACTGCTGA
GCTTACTGGAAAAGGACGAGTTCAAGTCTGGAGCCCCACGTGGACTTCTATGACAAAATAAGCCCATGTTTTTGGCT
TCCTGAACCTTTTTTGCCCCCACTTTTAGACATACCCCAGAGCCCTGTGGCTCCCCACACCCTGCCATAGGGGCAG
TCCTGCCTTACACATAGAAGCATTATGCCTGCTGCCCTGCCCTCAGGCACAGCCAGCTGTGAGCTCCCAGGTCA
TTGGCCTTACCAGTTGTCAGGAGTCTGTGCTGTGCACCCTGGGTTATAAGGAGGCTTAGGAGAGAGGTTATGGGT
ATTGGTGTCTCTATCCCCAGGAATAGAACTTAAGGGGTGGGAAGAACAGGAAAAGAAGCTGGAACACAGAAGAGA
GGAGGTTGTGCTCTTGCTCATAGCAAGCCTGTGGGTAGAGGAAAGAGTGATCTGGTGTGCAATAGGAGGACCCA
TGTAATTTCGAGATGGCCTGGATGGGAGGAAGGGCAGACGGTACATGTCCCAGCCCACATAGATGCCCTTGCT
GAGGGTAGCAGGACCTTCTGTTGAACTTTGTGTCCTCACTCTGATGTCTCCTTCCTTCAGAATCTACCACCCCTC
CCCCAGGCTGGGAGAAGGGGCTCCTGGGTGTCTGTATACACGCCAAAGGCAGATACAAATAAAATACAGATTGTC
CTTTAAAAAAAAAAAAAAAAAAAA

850/6881
FIGURE 790

MEGGLGRAVCLLTGASRGFGRTLAPLLASLLSPGSVLVLSARNDALRQLEAELGAERSGLRVVRVPADLGAEAG
LQQLLGALRELPRPKGLQRLLLINNAGSLGDIVSKGFVDLSDSTQVNNYWALNLTSMCLTSSVLKAFDPSGLNR
TVVNISSLALQPFKGWALYCAGKAARDMLFQVLALEEPNVRVLNYAPGPLDTDMQQLARETSVDPDMRKGLQEL
KAKGKLVDCVSAQKLLSLEKDEFKSGAHVDFYDK

851/6881
FIGURE 791

GGAGCATGGAGGGCGGGCTGGGGCGTGCTGTGTGCTTGCTGACCGGGGCTCCCGCGGCTTCGGCCGGACGCTGG
CCCCGCTCCTGGCCTCGCTGCTGTGCGCCGGCTCCGTGCTTGCTCCTTAGCGCCCGCAACGACGAGGCACTGCGCC
AGCTGGAGGCCGAGCTGGGCGCCGAGCGGTCTGGCCTGCGCGTGCTGCGGGTGCCCGCCGACCTGGGCGCCGAGG
CCGGCTTGCGAGCAGCTGCTCGGCGCCCTGCGCGAGCTCCCCCGGCCCAAGGGGCTGCAGCGACTGCTGCTTATCA
ACAACGCGGGTAAGACCCCGGGGCTGGAGCGGACTCCCCATGTGAGCGCCCACTTCCTCCACCGCTGGGGAATTT
AAGGGCTACTCCTAGGGGTCAGATAGGACGCCTAGAAATCTAAAGCCCTAGTTGTTCCAGCTTGAGTAGCAAGTG
GAGGCGAGGTGTACAGGAAGACTCGGTGCCGTGCGGACAACAGTCTGCCCCCACTTCCTGGAAAATTGGCCAGCC
GGGTTTTCCAAAGCGCCTGTAGAACTTTCCCGAGCCCTTGCTCTGCGCCCTCTGGCGGAGGCCTGAGTCAGGGC
TTGGAGTCTCCAAACAAGTAAACTCTCTGCTTCTCCAAAGGACCGCTAGAGCTTTCCTTCAGCCCCACGGCAAGG
GCTCTCTTGGGGATGTGTCCAAGGCTTCGTGGACCTGAGTGACTCCACTCAAGTGAACAATACTACTGGGCACTGA
ACTTGACCTCCATG

852/6881
FIGURE 792

GTTCCAAGGTTTTCGGGCCCGGTCTCGGAGAAGAGGGGAGAGTGGAGGGCCGCTGAATAAGCTTCCAAAATGATGC
CCACACCAGTTATCCTATTGAAAGAGGGGACTGATAGCTCCCAAGGCATCCCCAGCTTGTGAGTAACATCAGTG
CCTGCCAGGTGATTGCTGAGGCTGTAAGAACTACCTTGGGTCCCGTGGCATGGACAAGCTTATTGTAGATGGCA
GAGGCAAAGCAACAATTTCTAATGATGGGGCCACAATTCTGAAACTTCTTGATGTTGTCCATCCTGCAGCAAAGA
CTTTGGTAGACATTGCCAAATCCCAAGATGCTGAGGTGGGTGATGGCACCACCTCAGTGACCTTGGCTGGCTGCAG
AGTTTCTGAAGCAGGTGAAACCTATGTGGAGGAAGGTTTACACCCCGATCATCATTGAGCTTTCGCGCACAG
CCACCCAGCTGGCAGTTAACAAGATCAAAGAGATTGCTGTGACCGTGAAGAAGGCAGATAAAGTGGAGCAGAGGA
AGCTGCTGAAAAGTGTGCCATGACCGCTCTGAGCTCCAAGCTGATCTCCAGCAGAAAAGCTTTCTTTGCTAAGA
TGGTGGTGGATGCAGTGATGATGCTCGATGATTTGCTGCAGCTTAAAATGATTGGAATCAAGAAGGTACAGGGTG
GAGCCCTCGAGGATTCTCAGCTGGTAGCTGGTGTTCATTCAAGAAGACTTTCTCTTACGCTGGGTTTGAAATGC
AACCCAAAAGTACCACAATCCCAAGATTGCCCTTTTGAATGTCGAGCTCGAGTTGAAAGCTGAGAAAGACAATG
CTGAGATAAGAGTCCACACAGTTGAGGATTATCAGGCAATTGTTGATGCTGAGTGGAACATTCTCTATGACAAGT
TAGAGAAGATCCATCATTCTGGAGCCAAAGTTGTCTTGTCCAAACTCCCCATTGGGGATGTGGCCACCCAGTACT
TTGCTGACAGGGACATGTTCTGTGCTGGCCGAGTACCTGAGGAGGATCTGAAGAGGACAATGATGGCCTGTGGAG
GCTCAATCCAGACCAGTGTGAATGCTCTGTGTCAGCAGATGTGCTGGGTGATGCCAGGTGTTTGAAGAGACCCAGA
TTGGAGGCGAGAGGTACAATTTTTTACTGGCTGCCCAAGGCCAAGACATGCACCTTCATTCTCCGTGGCGGCG
CCGAGCAGTTTATGGAGGAGACAGAGCGGTCCCTGCATGATGCCATCATGATCGTCAGGAGGGCCATCAAGAATG
ATTGAGTGGTGGCTGGTGGCGGGGCCATTGAGATGGAACCTCTCCAAGTACCTGCGGGATTACTCAAGGACTATTC
CAGGAAAACAGCAGCTGTTGATTGGGGCTTATGCCAAGGCCTTGGAGATTATCCCACGCCAGCTGTGTGACAATG
CTGGCTTTGATGCCACAAACATTCTCAACAAGCTGCGGGCTCGGCATGCCAGGGGGGTACATGGTATGGAGTAG
ACATCAACAACGAGGACATTGCTGACAACCTTTGAAGCTTTCTGTGTTGGGAGCCAGCTATGGTGCAGTCAATGCCG
TGACAGCAGCCTCTGAGGCTGCGTGCCTGATCGTGTCTGTAGATGAAACCATCAAGAACCCCGCTCGACTGTGG
ATGCTCCCACAGCAGCAGGCCGGGGCCGTGGTCTGTGGCCGCCCCACTGAGAGGCACCCACCCATCACATGGCT
GGCTGGCTGCTGGGTGCACTTACCCTCCTTGGCTTGGTTACTTCATTTTACAAGGAAGGGGTAGTAATTGGCCCA
CTCTCTTCTTACTGGAGGCTATTTAAATAAAATGTAAGACTTCAAAAAAAAAAAAAAAAAAAAAA

853/6881

FIGURE 793

MMPTPVILLKEGTDSSQGIPQLVSNISACQVIAEAVRTTLGPRGMDKLIVDGRGKATISNDGATILKLLDVVHPA
AKTLVDIAKSQDAEVGDGTTSTLLAAEFLKQVKPYVEEGLHPQIIIRAFRTATQLAVNKIKEIAVTVKKADKVE
QRKLLKCAMTALSSKLISQQKAFFAKMVVDVAVMMLDDLLQLKMIGIKKVQGGALEDSQLVAGVAFKKTFSYAGF
EMQPKKYHNPKIALLNVELELKAEKDNAEIRVHTVEDYQAIVDAEWNILYDKLEKIHHSKAKVLSKLPIGDVAT
QYFADRMFCAGRVPEEDLKRTMMACGGSIQTSVNALSADVLGRCQVFEETQIGGERYNFFTGCPKAKTCTFILR
GGAEQFMEETERSLHDAIMIVRRAIKNDSVVAGGGAIEMELSKYLRDYSRTIPGKQQLLIGAYAKALEIIPRQLC
DNAGFDATNINLKLRRARHAQGGTWYGVDINNEDIADNFEAFVWEPAMVRINALTAASEAACLIVSVDETIKNPRS
TVDAPTAAGRGRGRGRPH

854/6881
FIGURE 794

ATATAACCGCGTGGCCCGCGCGCGCTTCCCTCCCGGCGCAGTCACCGGCGCGGTCTATGGCTGCGACTTCTCT
AATGTCTGCTTTGGCTGCCCCGGCTGCTGCAGCCCGCGCACAGCTGCTCCCTTCGCCTTCGCCCTTTCCACCTCGC
GGCAGTTTCGAAATGAAGCTGTTGTCAATTTCTGGAAGGAACTGGCCCAGCAGATCAAGCAGGAAGTGCGGCAGGA
GGTAGAAGAGTGGGTGGCCTCAGGCAACAAACGGGCCACACCTGAGTGTGATCCTGGTTGGCGAGAATCCTGCAAG
TCACTCCTATGTCCTCAACAAAACAGGGCAGCTGCAGTTGTGGGAATCAACAGTGAGACAATTATGAAACCAGC
TTCAATTTTCAGAGGAAGAATTGTTGAATTTAATCAATAAACTGAATAATGATGATAATGTAGATGGCCTCCTTGT
TCAGTTGCCTCTTCCAGAGCATATTGATGAGAGAAGGATCTGCAATGCTGTTTCTCCAGACAAGGATGTTGATGG
CTTTCATGTAATTAATGTAGGACGAATGTGTTTGGATCAGTATTCCATGTTACCGGCTACTCCATGGGGTGTGTG
GGAAATAATCAAGCGAACTGGCATTCCAACCCTAGGGAAGAATGTGGTTGTGGCTGGAAGGTCAAAAAACGTTGG
AATGCCCATTGCAATGTTACTGCACACAGATGGGGCCGATGAACGTCCCGGAGGTGATGCCACTGTTACAATATC
TCATCGATAATACTCCCAAAGAGCAGTTGAAGAAACATACAATTCCTTGCAATATTGTAATATCTGCTGCAGGTAT
TCCAAATCTGATCACAGCAGATATGATCAAGGAAGGAGCAGCAGTCATTGATGTGGGAATAAATAGAGTTCCGA
TCCTGTAACAGTCCAAACCCAGTTGGTTGGAGATGTGGATTTTGAAGGAGTCAGACAAAAGCTGGGTATATCAC
TCCAGTTCCCTGGAGGTGTTGGCCCCATGACAGTGGCAATGCTAATGAAGAATACCATTATTGCTGCAAAAAAGGT
GCTGAGGCTTGAAGAGCGAGAAGTGCTGAAGTCTAAAGAGCTTGGGGTAGCCACTAATTAACTACTGTGCTTCT
GTGTCACAAACAGCACTCCAGGCCAGCTCAAGAAGCAAAGCAGGCCAATAGAAATGCAATATTTTAAATTTATTC
TACTGAAATGGTTTAAAAATGATGCCTTGTATTTATTGAAAGCTTAAATGGGTGGGTGTTTCTGCACATACCTCTG
CAGTACCTCACCAGGGAGCATTCCAGTATCATGCAGGGTCTGTGATCTAGCCAGGAGCAGCCATTAACCTAGTG
ATTAATATGGGAGACATTACCATATGGAGGATGGATGCTTCACTTTGTCAAGCACCTCAGTTACACATTTCGCCTT
TTCTAGGATTGCATTTCCCAAGTGCTATTGCAATAACAGTTGATACTCATTTTAGGTACCAGACCTTTTGAGTTC
AACTGATCAAACCAAAGGAAAAGTGTGCTAGAGAAAATTGGGGAAAAGGTGAAAAAGAAAAATGGTAGTAATT
GAGCAGAAAAAATTAATTTATATATGATTGATTGGCAACCAGATTTATCTAAGTAGAACTGAATTGGCTAGGA
AAAAAGAAAACTGCATGTTAATCATTTTCCTAAGCTGTCCTTTTGAGGCTTAGTCAGTTTATTGGGAAAATGTT
TAGGATTATTCCTTGCTATTAGTACTCATTTTATGTATGTTACCCTTCAGTAAGTTCTCCCCATTTTAGTTTTCT
AGGACTGAAAGGATTCTTTTCTACATTATACATGTGTGTTGTCATATTTGGCTTTTGCTATATACTTTAACTTCA
TTGTTAAATTTTGTATTGTATAGTTTCTTTGGTGTATCTTAAACCTATTTTGAAAAACAACTTGGCTTGAT
AATCATTTGGGCAGCTTGGGTAAGTACGCAACTTACTTTTCCACCAAAGAACTGTCAGCAGCTGCCTGCTTTTCT
GTGATGTATGTATCCTGTTGACTTTTCCAGAAATTTTTAAGAGTTTGAGTTACTATTGAATTTAATCAGACTTT
CTGATTAAAGGGTTTTCTTTCTTTTTTAATAAAACACATCTGCTCTGGTATGGTA

855/6881
FIGURE 795

MSALAARLLQPAHSCSLRLRPFHLAAVRNEAVVISGRKLAQQIKQEVQRQVEEWVASGNKRPHLSVILVGENPAS
HSYVLNKTRAAAVVGINSETIMKPASISSEELLNLINKLNDDNVDGLLVQLPLPEHIDERRICNAVSPDKDVDG
FHVINVGRMCLDQYSMLPATPWGVWEIIKRTGIPTLGKNVVVAGRSKNVGMPIAMLLHTDGAHERPGGDATVTIS
HRYTPKEQLKKHTILADIVISAAGIPNLITADMIKEGAAVIDVGINRVHDPVTAKPKLVGDVDFEGVRQKAGYIT
PVPGGVGPMTVAMLMKNTIIAAKKVLRLEEREVLKSKELGVATN

856/6881
FIGURE 796

GGAAAAATTGTACAGCAGCCAGAGGGGTTTAAACAGGAGTGCAGAGGGATAAGGGCAGCTTCTGCCCTCTGCCCCA
AGAGCTGGCCACCTCTTTAAAGACTGAGGGAACAGTGGGAGGAGGAACCTGTGGGACAGTGTGGTACCTATCTGTG
CCCCCTCTGGAGGGGTTGACAAGGGAAAGGGCACCGGGGGGCACAGAGATGCAGGACAGATTGCACATCCTGGAG
GACCTGAATATGCTCTACATTCGGCAGATGGCACTCAGCCTGGAGGACACGGAGTTGCAGAGGAAGCTAGACCAT
GAGATCCGATGAGGGAAGGGGCTGTAAAGCTGCTGGCAGCCTGCTCCCAGCGAGAGCAGGCTCTGGAGGCCACC
AAGAGCCTGCTAGTGTGCAACAGCCGCAATCCTCAGCTACATGGGCGAGCTGCAGCGGCGCAAGGAGGCGCAGGTG
CTGGGGAAGACAAGCCGGCGGCTTCTGACAGTGGCCCGCCGCTGAGCGCTCCCCCTGCCGCGGCGGGTCTGTC
ATCTCTGACCTCCGGATTCCACTCATGTGGAAGGACACAGAATATTTCAAGAACAAAGGTGACTTGACCCGCTGG
GCTGTGTTCTGTGCTGCTGCACTGGGGGAACACATCCAGGACACAGAGATGATCCTAGTGGACAGGACCCTCACA
GACATCTCCTTTAGAGCAATGTGCTCTTCGCTGAGGCGGGGCCAGACTTTGAAGTGCAGGTTAGAGCTGTATGGG
GCCTGTGTGGAAGAAGAGGGGGCCCTGACTGGCGGGCCCAAGAGGCTTGCCACCAAAGTCTCAGCAGTCCCTGGGC
CGCTCCTCAGGGAGGCGTGTCCGGGCATCGCTGGCAGTGTGGGGTTTCAGGGAGCAGTCCCATCTTGTCTCCCC
ACCCAGTTGTTGGTGGTCCCTCGTTACCACCTCTTGGCTCACACCACACTCACCTGGCAGCAGTGCAAGATGGA
TTCCGCACACATGACCTCACCTTGCCAGTCATGAGGAGAACCCTGCCTGGCTGCCCTTTATGGTAGCGTGTGT
TGCCGTCTGGCAGCTCAGCCTCTCTGCATGACTCAGCCACTGCAAGTGGTACCTCAGGGTGCAGCAAGCTGGG
GAGATGCAGAACTGGGCACAAGTGCATGGAGTTCTGAAAGGCACAAACCTCTTCTGTTACCGGCAACCTGAGGAT
GCAGACACTGGGGAAGAGCCGCTGCTTACTATTGCTGTCAACAAGGAGACTCGAGTCCGGGCAGGGGAGCTGGAC
CAGGCTCTAGGACGGCCCTTCACCCTAAGCATCAGTAACCAGTATGGGGATGATGAGGTGACACACACCCTTCAG
ACAGAAAGTCGGGAAGCACTGCAGAGCTGGATGGAGGCTCTGTGGCAGCTTTTCTTTGACATGAGCCAATGGAAG
CAGTGTGTGATGAAATCATGAAAATTGAACTCCTGCTCCCCGGAAACCACCCCAAGCACTGGCAAAGCAGGGG
TCCTTGATACATGAGATGGCTATTGAGCCGCTGGATGACATCGCAGCGGTGACAGACATCCTGACCCAGCGGGAG
GGCGCAAGGCTGGAGACACCCCAACCTGGCTGGCAATGTTTACAGACCAGCCTGCCCTGCCTAACCCCTGCTCG
CCTGCCTCAGTGGCCCCAGCCCCAGACTGGACCCACCCCTGCCCTGGGGGAGACCCCGAACCTTTTCCCTGGAT
GCTGTCCCCCAGACCACTCCCCTAGGGCTCGCTCGGTTGCCCCCTCCCACCTCAGCGATCCCCACGGACCAGA
GGCCTCTGCAGCAAAGGCCAACCTCGCACTTGGCTCCAGTCACCAGTGTGAGAGAGAAAGGTGCTGGCATAGGAT
CTGCCCAGAAGAGAAAATGACCCATGCGCAGTTGGGCTCTGGATACGGCGCTGTCTATAGCAAGTTGGCCAGTCT
GGCCTCCTGTTTCTCTGCTGGACCTGGGGTAGGCTGCAGGGGTGGGCAGAAGCCCCTCTTAAATTGTGGTTGCCA
TGGTACCGAGGGACTCATTCTGGGGCTCGCTGGGACCTCCCTAAACCCTTCCTGGAAGAAAAGTGAACCAACT
CTGCCCTACCTCCCTGCACTAACCAGCTTTGAGGATGGCACTGAAGAACCCTTGGAGCAAACATACCTCCCTTGT
GACTCCACATCAACCATTAAAGTTATTTAACAGCAAAAAAAAAAAAAA

857/6881
FIGURE 797

MQDRLHILEDLNMLYIROMALSLEDTELQRKLDHEIRMREGACKLLAACSQREQALEATKSLVCNSRILSYMGE
LQRRKEAQVLGKTSRRPSDSGPPAERSPCRGRVCISDLRIPLMWKDTEYFKNKGDHLRWAVFLLLQLGEHIQDTE
MILVDRTLTDISFQSNVLF AEAGPDFELRLELYGACVEEEGALTGGPKRLATKLSSSLGRSSGRRVRASLDSAGG
SGSSPILLPTPVVGGPRYHLLAHTTLTAAVQDGFRTHTLTLASHEENPAWLPLYGSVCCRLAAQPLCMTQPTAS
GTLRVQQAGEMQNWAQVHGVLKGTNLFCYRQPEDADTGEEPLLTIAVNKETRV RAGELDQALGRPFTLSISNQYG
DDEVTHLTQTESREALQSWMEALWQLFFDMSQWKQCCDEIMKIETPAPRKPPQALAKQGSLYHEMAIEPLDDIAA
VTDILTQREGARLET PPPWLAMFTDQPALPNPCSPASVAPAPDWITHPLPWGRPRTFSLDAVPPDHSRARSVAPL
PPQRSPTRGLCSKGQPRTWLQSPV

858/6881
FIGURE 798

GACTACGTGGGTCITGGAGCTGACTGCCGTCCTGACACGTCCTAGAGCTGCAAGTCCTGCCAGAGAGCCACCATGA
CCTCTCAGCCTCTCAGGCTAGCAGAAGAGTATGGCCCAAGTCCTGGGGAGTCTGAACCTGGCTGTGAACCCCTTTG
ATGGGCTTCCCTTCTCTTCCCGCTACTATGAGCTGCTGAAGCAGCGCCAAGCCTTGCCCATCTGGGCTGCTCGCT
TTACCTTCTTGGAGCAGTTGGAGAGTAACCCCACTGGAGTGGTGTCTGGTGTCTGGGGAGCCTGGTTCTGGCAAGA
GCACCCAGATCCCTCAGTGGTGTGCAGAGTTTGCCTGGCCAGAGGGTTCAGAAAGGACAGGTTACTGTTACTC
AGCCCTACCCTCTTGACGCCCGGGAGCCTGGCTCTGCGGGTTGCTGATGAGATGGACCTGACCCTGGGTCATGAG
GTTGGATCCAGCATCCCCAGGAGGACTGCACGGGGCCCAACACCTGCTCAGGTTCTGCTGGGACAGGCTGCTT
CTGCAGGAGGTGGCCTCGACCCGAGGCACTGGAGCCTGGGGCGTGTGGTACTAGATGAGGCTCAGGAGCGGTGCG
GTGGCATCAGATTCACTCCAGGGGCTACTGCAAGATGCCAGGCTGGAAAACTTCCGGGGGACCTCAGAGTGGTT
GTGGTTACTGACCCAGCCCTTGAACTAAGCTCCGAGCTTCTGGGGCAATCCTCCTATTGTGCATATACCCAGA
GAGCCTGGTGAGAGACCTTCCCCATCTACTGGGACACCATCCACCTGATCGGGTGGAAGCTGCCTGCCAAGCA
GTGCTTGAATTGTGTGCGAAGGAGCTTCCAGGAGATGTGCTAGTGTCTGCTGCCAGTGAGGAGGAAATTTCCCTG
TGCTGTGAATCCTTGTCAGGGAGGTAGAGTCCTTGCTTCTCCAAGGGCTTCCACCAGAGTACTGCCCTTAC
CCAGACTGTGGACGAGCCGTTCCAGCTGTGTATGAGGACATGGATGCCGAAAGGTTGTGGTCACTCACTGGCTG
GCTGACTTCTCCTTCTCCCTCCCTTCCATCCAACATGTGCTGACTCAGGACTGGAGCTCCGAAGTGTTTACAA
CCTAGGATCCGAGCAGAATTCCAAGTGTGAGGCCAATCAGCAAGTGTGAGGCAGAGGCAAGACGATTGCCAGCA
AGAGGGTTCCACAGGATCCTGCCTCTGCCTGTATCCTAAGTCCTTCTAGAACTAGAAGCTCCACCATTGCCA
CAACCCAGGGTGTGTGAGGAGAATCTGAGCTCCCTGGTGTACTACTAAAAAGGAGACAGATTGCAGAGCCAGGG
GAGTGTCACTTCTGGACAGCCTGCTCCAGAAGCACTGATGCAAGCCCTGGAAGATTTAGACTATCTGGCAGCC
CTGGATGATGATGGGGACCTGTGAGATCTGGGTGTCTACTATCAGAATTCCTCTGGCCCTGAGCTGGCCAAA
GCCCTGCTGGCCTCATGCGAGTTTGAAGTGTGTGGACGAGATGCTCACCCTGGCTGCCATGCTCACAGCTGCCCT
GGGTTTACCCGTCCTCCACTCAGTGCAGAAGAAGCTGCCCTGCGTCGGGCCCTGGAACACACGGATGGTGACCAC
AGTTCTCTGATCCAGGTGTATGAAGCCTTTATACAAAGTGGAGCAGATGAGGCTTGGTGCCAGGCTCGAGGTCTG
AATTGGGCAGCATTGTGCCAAGCCCATAACTTCCGGGGAGAATCCTAGAACTCATGCAACGAATTGAACCTCCC
TTGTCCCTACCAGCCTTTGGCTCTGAGCAGAATCGCAGAGACCTTCAGAAAGCACTGGTGTGAGAATACTTTCTC
AAGGTGGCCAGAGACACAGACGGGACTGGAAATTACCTTCTCCTAACCATAAGCATGTGGCCAGCTCTCCTCA
TACTGCTGCTACCGAAGCCGCAGAGCTCCTGCCAGACCCCCACCATGGGTGCTCTACCACAATTTACCATATCC
AAAGACAACCTGCCTCTCCATTGTTTCTGAGATTCAACCACAGATGCTGGTGGAATTGGCCCTCCATACTTCTG
AGTAACTTGCCCTCCAGTGAGAGCAGAGACCTTCTGAACCAGCTAAGGGAAGGAATGGCAGATTCTACAGCAGG
AGCAAATCATCTCAGCCCAGGAGTTCAGAGATCCCTGTGTCCTGCAGTGGACCTGCCTGCCTATGGAATGGAGCT
GGGTTTCATCTCATCACATTAGATTATCCCTCAGGGTGACACCAAGCACCCAGACAGATTTAGAAGCCCAAAGTT
TAGGGTCAAATGTAAACCTGGAACCTGAGTCCCAAGAAATGGTAGACTGGGAATGGAAGAATGGGGTAAACCA
CAGTCTACATAGGGAAGGACTCTTTCCTTAGCCTTCTCTTATTGATTGGAGAGGGACTGACATGCTCCTCATTCT
CTTAACCTTGCCAAACCCATTCTTGTAATCCTTGTGATCTATAAAAGATTTTCTATGATGCCAAAAA
AAAAA

859/6881
FIGURE 799

MDLTILGHEVGSSIPQEDCTGPNITLLRFCWDRLLLQEVASTRGTGAWGVVLVDEAQERSVASDSLQGLLDARLEK
LPGDLRVVVVTDPALEPKLRAFWGNPPIVHIPREPGERPSPYWDITPPDRVEAACQAVLELCRKELPGDVLVFL
PSEEEISLCCESLSREVESLLLQGLPPRVLPPLHPDCGRAVQAVYEDMDARKVVVTHWLADFSFSLPSIQHVIDSG
LELRSVYNPRIRAEFQVLRPISKQAEARRLRARGFPPGSCCLYPKSFLELEAPPLPQPRVCEENLSSLVLLK
RRQIAEPGECHFLDQPAPEALMQALEDLDYLAALDDGDLSDLGVILSEFPLAPELAKALLASCEFCVDEMLTL
AAMLTAAPGFTRPPLSAEEAALRRALEHTDGDHSSLIQVYEAFIQSGADEAWCQARGLNWAALCQAHKLRGELLE
LMQRIELPLSLPAFGSEQNRRDLQKALVSEYFLKVARDTDGTGNYLLLTHKHVAQLSSYCCYRSRRAPARPPWV
LYHNFTISKDNCLSIQVSEIQPQMLVELAPPYFLSNLPPSESRLNQLREGMADSTAGSKSSSAQEFRDPCVLQ

860/6881
FIGURE 800

CGAAAAATGGCGGCGGCGGCGACGGCCGGGCGCTCCTGAAGCAGCAGTTATGGAGCTTCCCTCAGGGCCGGGGCCG
GAGCGGCTCTTTGACTCGCACCGGCTTCCGGGTGACTGCTTCCTACTGCTCGTGCTGCTGCTCTACGCGCCAGTC
GGGTTCTGCCTCCTCGTCTGCGCCTCTTTCTCGGGATCCACGTCTTCCTGGTCAGCTGCGCGCTGCCAGACAGC
GTCCTTCGCAGATTCTAGTAGCGGACCATGTGTGCGGTGCTAGGGCTCGTGGCCCCGGCAGGAGGACTCCGGACTC
CGGGATCAGAGTGTGAGGGTCCCTCATTTCACACCATGTGACACCTTTTCGACCACACATAGTCAATTTGCTTACC
ACCTGTAGCACCCCTCTACTCAATAGTCCCCCAGCTTTGTGTGCTGGTCTCGGGGCTTCATGGAGATGAATGGG
CGGGGGGAGTTGGTGGAGTCACTCAAGAGATTCTGTGCTTCCACGAGGCTTCCCCCACTCCTCTGCTGCTATTCT
CCTGAGGAAGAGGCCACCAATGGCCGGGAGGGGCTCCTGCGCTTCAGTTCCCTGGCCATTTTCTATCCAAGATGTG
GTACAACCTCTTACCCTGCAAGTTCAGAGACCCCTGGTCTCTGTGACGGTGTGAGATGCCTCCTGGGTCTCAGAA
CTGCTGTGGTCACTTTTTCTCCCTTTTACGGTGTATCAAGTAAGGTGGCTTCGTCTGTTTCATCGCCAAGTAGGG
GAAGCGAATGAGGAGTTTGCACTCCGTGTACAACAGCTGGTGGCCAAGGAATTGGGCCAGACAGGGACACGGCTC
ACTCCAGCTGACAAAGCAGAGCACATGAAGCGACAAAGACACCCAGATTGCGCCCCAGTCAGCCCAGTCTTCT
TTCCCTCCCTCCCTGGTCTTCTCCTGATGTGCAACTGGCAACTCTGGCTCAGAGAGTCAAGGAAGTTTGGCC
CATGTGCCATTGGGTGTCTATCCAGAGAGACCTGGCCAAGACTGGCTGTGTAGACTTGACTATCACTAATCTGCTT
GAGGGGGCCGTAGCTTTTCATGCCTGAAGACATACCAAGGGAAGTCACTCCCTACCCACAGCCTCTGCCTCCAAG
TTTCCCAGCTCTGGCCCCGTGACCCCTCAGCCAACAGCCCTAACATTTGCCAAGTCTTCCTGGGCCCGGCAGGAG
AGCCTGCAGGAGCGCAAGCAAGCACTATATGAATACGCAAGAAGGAGATTACAGAGAGACGAGCCCAGGAGGCT
GACTGAGCTCAAAGGAACAGGATGGCACCCAGAGCCGAGGAGACTGGGGGCAGCCCTACCCAACTCACA
ACAGGCTGGATGGGTGGGTGGTAAAAAGGAAGGATGAGGCTCCCCCAATGTCACATTAAATTCATGGTTTTCAT
TC

861/6881
FIGURE 801

CGTGGATCCCGAGAAAGAGGCGCAGGACGAGGAGGCAGAACCCGACTGGCGCGTAGAGCAGCAGCACGAGCAGTA
GGAAGCAGTCACCCGGAAGCCTGGGGGCGAGAGGCGAAGTGGTCAGGCGCCGAAGGCCGAGAGCACGCGGGGATC
GGTCTCTTCCCGCCGGGTCTCTTACCGGTGCGAGTCAAAGAGCCGCTCCGGCCCCGGCCCTGAGGGAAGCTCCAT
AACTGCTGCTTCAGGAGCGCCCGGCCGTCGCCGCCGCCGCAATTTTCGCGCCCGGCCGAGGGGCTCTTGGGAAG
GCGGAGTCTTTGGGCATCCGCCCGGGGTGAGGGGACCCGAAGTCTTGAGGCGCGCCGAAGGGCTAGCGGTCCCA
GCATACCCCGCGGCCCTTGGGCCGTCTCACAACCTCGCGTCCGGCGGAGACCACAATTCCCGGCATTCTGTGGGGC
AGGGAGGAGTCGGCCTCCCGGAATCCTGGTCCCGGCGTGCACTTCTGAAGGACTTCAGGTACCGCGGTGCCCCGC
GTCCTACTGTCCGCCTGCTCGCGTCTGGGTGCCGCTCTGAGTAGGGCGGGCGAGGAGGCAGCCAAGGCGGAGC
TGATGGCTGCGCCGAGGGCGGGGCGGGGTGCAAGGCTGGAGCCTTCGGGCATGGCGGGCTTTGGGGGGCATTCTGCT
GGGGGAGGAGACCCCGTTTGACCCCTGACCTCCGGGCCCTGCTGACGTCAGGAACCTCTGACCCCGGGCCCGAG
TGACTTATGGGACCCCCAGTCTCTGGGCCCGGTTGTCTGTTGGGGTCACTGAACCCGAGCATGCCTGACGCTCTG
GGACCCCGGGTCCCCGGGCACAACCTGACTGCGGTGACCCAGATACCAGGACCCGGGAGGCCTCAGAGAACTCTG
GAACCCGTTTCGCGCGCGTGCTGGTGGCGGTGGCGCTGGGCGCTGGGGGGCAGTGCTGTGTTGTTGTTGGGGCGGG
GTCGGGGTCTCTCCGGCCGTCCTCGCCGCCGTCCTTAGCCCGCCGCCCGCTTCTCCCCGGAGTCAGTACAACCTCA
TCGAGATGTGGTGGAGAAGACAGCACCTGCCGTGGTCTATATCGAGATCCTGGACCGGCACCCTTTCTTGGGCC
GCGAGGTCCTATCTCGAACGGCTCAGGATTCTGGTGGCTGCCGATGGGCTCATTGTACCAACGCCCATGTGTG
TGGCTGATCGGCGCAGAGTCCGTGTGAGACTGCTAAGCGGCGACACGTATGAGGCCGTGGTCACAGCTGTGGATC
CCGTGGCAGACATCGCAACGCTGAGGATTCTAGACTAAGGAGCCTCTCCCCACGCTGCCTCTGGGACGCTCAGCTG
ATGTCCGGCAAGGGGAGTTTGTGTTGCCATGGGAAGTCCCTTTGCACTGCAGAACACGATCACATCCGGCATTG
TTAGCTCTGCTCAGCGTCCAGCCAGAGACCTGGGACTCCCCAAACCAATGTGGAATACATTCAAACCTGATGCAG
CTATTGATTTTGGAAACTCTGGAGGTCCCCGTGGTTAACCTGGATGGGAGGTGATTGGAGTGAACACCATGAAGG
TCACAGCTGGAATCTCCTTTGCCATCCCTTCTGATCGTCTTCGAGAGTTTCTGCATCGTGGGGAAAAGAAGATT
CCTCCTCCGGAATCAGTGGGTCCCAGCGGCGCTACATTGGGGTGATGATGCTGACCCTGAGTCCCAGCATCCTTG
CTGAACTACAGCTTCGAGAACCAAGCTTTCCCGATGTTTCAGCATGGTGTACTCATCCATAAAGTCATCCTGGGCT
CCCCCTGCACACCGGGCTGGTCTGCGGCCTGGTGATGTGATTTTGGCCATTGGGGAGCAGATGGTACAAAATGCTG
AAGATGTTTATGAAGCTGTTTCGAACCAATCCAGTTGGCAGTGCAGATCCGGCGGGGACGAGAAACACTGACCT
TATATGTGACCCCTGAGGTACAGAAATGAATAGATACCAAGAGTATGAGGCTCCTGCTCTGATTTCCTCCTTGC
CTTTCTGGCTGAGGTTCTGAGGGCACCGAGACAGAGGGTTAAATGAACCAAGTGGGGCAGGTCCCTCCAACCACC
AGCACTGACTCCTGGGCTCTGAAGAATCACAGAAACACTTTTTATATAAAATAAAATTATACCTAGCAACATATT
ATAGTAAAAAATGAGGTGGGAGGGCTGGATCTTTTCCCCCACCAAAAGGCTAGAGGTAAAGCTGTATCCCCCTAA
ACTTAGGGGAGATACTGGAGCTGACCATCCTGACCTCCTATTAAAGAAAATGAGCTGCTGAAAAAAAAAAAAAA
A

862/6881
FIGURE 802

MAAPRAGRAGWSLRAWRALGGIRWGRRPRLTPDLRALLTSGTSDPRARVTYGTPSLWARLSVGVTEPRACLTSG
TPGPRAQLTAVTPDTRTREASENSGTRSRAWLAVALGAGGAVLLLLWGGGRGPPAVLAAVSPPPASPRSQYNFI
ADVVEKTAPAVVYIEILDRHPFLGREVPISNGSGFVVAADGLIVTNAHVADRRRVVRLLSGDTYEAVVTAVDP
VADIATLRIQTKEPLPTLPLGRSADVRQGEFVVMGSPFALQNTITSGIVSSAQRPARDLGLPQTNVEYIQTDA
IDFGNSGGPLVNLDGEVIGVNTMKVTAGISFAIPSDRLREFLHRGEKKNSSSGISGSQRRYIGVMMLTTLSPSILA
ELQLREPSFPDVQHGVLIHKVILGSPAHRAGLRPGDVILAIGEQMVQNAEDVYEAVRTQSQLAVQIRRGRETLL
YVTPEVTE

863/6881

FIGURE 803

GCAACGCGAGTGGGAGCACCAGGATCTCGGGCTCGGAACGAGACTGCACGGATTGTTTTAAGAAAATGGCAGACA
AACCAGACATGGGGGAAATCGCCAGCTTCGATAAGGCCAAGCTGAAGAAAACGGAGACGCAGGAGAAGAACACCC
TGCCGACCAAAGAGACCATTGAGCAGGAGAAGCGGAGTGAAATTTCCCTAAGATCCTGGAGGATTCCTACCCCG
TCCTCTTCGAGACCCAGTCGTGATGTGGAGGAAGAGCCACCTGCAAGATGGACACGAGCCACAAGCTGCACTGT
GAACCTGGGCACTCCGCGCCGATGCCACCGGCCTGTGGGTCTCTGAAGGGACCCCCCAATCGGACTGCCAAA
TTCTCCGGTTTGCCCCGGGATATTATAGAAAATTATTTGTATGAATAATGAAAATAAAACACACCTCGTGGCATG
GCA

864/6881
FIGURE 804

MADKPDMGGEIASFDKAKLKKTTETQEKNILPTKETIEQEKRSIS

865/6881
FIGURE 805

CACCAAATGGCGGATGACGCCGGTGCAGCGGGGGGGCCAGAGCCCTGGTGGCCATGGGATAGGGAACCGCGGTG
GCTTCCACGGAGGTTTCGGCAGTGGCATCCGGGGCCGGGGTCGCGGGAGTGGACGGGGCCGTGCCGAGGCCGCGG
AGCTCGCGGAGGCAAGGCCGAGGATAAGGAGTGGATGCCTGTCACCAAGCTGGGCGCTTGGTCAAGGACATGAA
GATCAAGTCCCTGGAGGAGATCTATCTCTTCTCCCTGCCCATTAAAGGAATCTGAGATCATTGATTCTTCTCTGGG
GGACTCTCTCAAGGATGAGGTTTTGAAGATTATGCCGGTGCAGAAGCAGACCCGTGCCGGCCAGCGCACCAGGTT
CAAGGTGACAGGCCGCTGCGGCTCTGTGCTGGTGCGCCTCATCCCTGCACCCAGGGGCACTGGCATCGTCTCCGC
ACCTGTGCCCCAAGAAGCTGCTCATGATGGCTGGTATCGATGACTGCTACACCTCAGCCCCGGGGCTGCACTGCCAC
CCTGGGCAACTTCGCCAAGGCCACCTTTGATGCCATCTCTAAGACCTACAGCTACCTGACCCCCGACCTCTGGAA
GGAGACTGTATTTACCAAGTCTCCCTATCAGGAATCACTGACCACCTCGTCAAGACCCACACCAGAGTCTCGGT
GCAGCGGACCCAGGCTTCAGATGTGGCTACAACATAGGGTTTTTATACAAGAAAAATAAAGTGAATTAAGCCTG

866/6881
FIGURE 806

CGCAGGCTGGAAGGAAGACGAACCTACGAAGCAGAGATCTGAAGACAGCATGTACACAGCCATTCCCCAGAGTGG
CTCTCCATTCCCAGGCTCAGTGCAGGATCCAGGCCTGCATGTGTGGCGGGTGGAGAAGCTGAAGCCGGTGCCTGT
GGCGCAAGAGAACCAGGGCGTCTTCTTCTCGGGGGACTCCTACCTAGTGTGCACAATGGCCCAGAAGAGGTTTC
CCATCTGCACCTGTGGATAGGCCAGCAGTCATCCCGGGATGAGCAGGGGGCCTGTGCCGTGCTGGCTGTGCACCT
CAACACGCTGCTGGGAGAGCGGCCTGTGCAGCACCGCGAGGTGCAGGGCAATGAGTCTGACCTCTTCATGAGCTA
CTTCCCACGGGGCCTCAAGTACCAGGAAGGTGGTGTGGAGTCAGCATTTACAAGACCTCCACAGGAGCCCCAGC
TGCCATCAAGAACTCTACCAGGTGAAGGGGAAGAAGAACATCCGTGCCACCGAGCGGGCACTGAAGTGGGACAG
CTTCAACACTGGGGACTGCTTCATCCTGGACCTGGGCCAGAACATCTTCGCTGGTGTGGTGGAAAGTCCAACAT
CCTGGAACGCAACAAGGCGAGGGACCTGGCCCTGGCCATCCGGGACAGTGAGCGACAGGGCAAGGCCAGGTGGA
GATTGTCACTGATGGGGAGGAGCCTGCTGAGATGATCCAGGTCTGGGCCCCAAGCCTGCTCTGAAGGAGGGCAA
CCCTGAGGAAGACCTCACAGCTGACAAGGCAAATGCCCAGGCCGAGCTCTGTATAAGGTCTCTGATGCCACTGG
ACAGATGAACCTGACCAAGGTGGCTGACTCCAGCCCCTTTGGCCTTGAAGTGTGATATCTGATGACTGCTTTGT
GCTGGACAACGGGCTCTGTGGCAAGATCTATATCTGGAAGGGGCGAAAAGCGAATGAGAAGGAGCGGCAGGCAGC
CCTGCAGGTGGCCGAGGGCTTCATCTCGCGCATGCAGTACGCCCCGAACACTCAGGTGGAGATTCTGCCTCAGGG
CCGTGAGAGTCCCATCTTCAAGCAATTTTCAAGGACTGGAATTCAGGGGTGGGCGTCTTCTGCCCCATGCTCCC
CTGCCCCCACCACCTGCCTGCTTGCTTCTCTGGCTGCCTGGTCAGTGCAGAGGTGCCCCCTGCAGATGTTCAAT
AAAGGAGACAAGTGCTTTCCC

867/6881
FIGURE 807

MYTAIPQSGSPFPQSVQDPGLHVWRVEKLPVPVAQENQGVFFSGDSYLVLHNGPEEVSHLHLWIGQQSSRDEQG
ACAVLAVHLNTLLGERPVQHREVQGNESDLFMSYFPRGLKYQEGGVESAFHKTSTGAPAAIKKLYQVGKKKNIRA
TERALNWDSFNTGDCFILDGQNIFAWCGGKSNILERNKARDLALAIRDSERQGAQVEIVTDGEEPAEMIQVLG
PKPALKEGNPEEDLTADKANAAALYKVSDATGQMNLTQVADSSPFALELLISDDCFVLDNGLCGKIYIWKGRK
ANEKERQAALQVAEGFISRMQYAPNTQVEILPQGRESPIFKQFFKDWK

868/6881
FIGURE 808

GGCACGAGGCGCCCGCCTGCTACGAGTAGAACGCTGTCCGCAGCTTGCGCATTTGCGAGCCGCTGCCGCTCGCC
GCTGCTCCTTCGTAAGGCCACTTCCGCACACCGACACCAACATGAACGGACAGCTCAACGGCTTCCACGAGGCGT
TCATCGAGGAGGGCACATTCTCTTTTACCTCAGAGTCGGTCGGGGAAGGCCACCCAGATAAGATTTGTGACCAAA
TCAGTGATGCTGTCTTGTATGCCCACCTTCAGCAGGATCCTGATGCCAAAGTAGCTTGTGAACTGTTGCTAAAA
CTGGAATGATCCTTCTTGTGCTGGGGAATTACATCCAGAGCTGCTGTTGACTACCAGAAAGTGGTTCTGTAAGCTG
TTAAACACATTGGATATGATGATTCTTCCAAAGGTTTTGACTACAAGACTTGTAACGTGCTGGTAGCCTTGGAGC
AACAGTCACCAGATATTGCTCAAGGTGTTTCATCTTGACAGAAATGAAGAAGACATTGGTGCTGGAGACCAGGGCT
TAATGTTTGGCTATGCCACTGATGAAACTGAGGAGTGATGCCCTTAACCATTGTCTTGGCACACAAGCTAAATG
CCAACTGGCAGAACTACGCCGTAATGGCACTTTGCCTTGGTTACGCCCTGATTCTAAACTCAAGTTACTGTGC
AGTATATGCAAGGATCGAGGTGCTGTGCTTCCCATCAGAGTCCACACAATTGTTATATCTGTTTCAGCATGATGAAG
AGGTTTGTCTTGATGAAATGAGGGATGCCCTAAAGGAGAAAGTCATCAAAGCAGTTGTGCCTGCGAAATACCTTG
ATGAGGATACAATCTACCACCTACAGCCAAGTGGCAGATTTGTTATTGGTGGGCCTCAGGGTGATGCTGGTTTGA
CTGGACGCAAAATCATTGTGGACACTTATGGCGGTTGGGGTGCTCATGGAGGAGGTGCCTTTTCAGGAAAGGATT
ATACCAAGGTCGACCGTTCAGCTGCTTATGCTGCTCGTTGGGTGGCAAAATCCCTTGTTAAAGGAGGTCTGTGCC
GGAGGGTTCTTGTTCAGGTCTCTTATGCTATTGGAGTTTCTCATCCATTATCTATCTCCATTTTCCATTATGGTA
CCTCTCAGAAGAGTGAGAGAGAGCTATTAGAGATTGTGAAGAAGAATTCGATCTCCGCCCTGGGGTCAATGTCA
GGGATCTGGATCTGAAGAAGCCAATTTATCAGAGGACTGCAGCCTATGGCCACTTTGGTAGGGACAGCTTCCCAT
GGGAAGTGCCCAAAAGCTTAAATATGAAGTGTTAGCCTTTTTTCCCCAGACTTGTGGCGTAGGCTACAGAG
AAGCCTTCAAGCTCTGAGGGAAGGGCCCTCCTTCCATAAATTTTCTGTCTCTTTCAGCTCCTGACCAGTTGCA
GTCACCTCTAGTCAATGACATGAATTTTAGCTTTTGTGGGGGACTGTAAGTTGGGCTTGCTATTCTGTCCCTAGGT
GTTTTGTTACCATTATAATGAATTTAGTGAGCATAGGTGATCCATGTAAGTGCCTAGAAACAACACTGTAGTAA
ATAATGCTTTGAAATTGAACCTTTGTGCCCTATCACCAACGCTCCAAAGTCATAATTGCATTGACTTTCCCCAC
CAGATGCTGAAATGTCTTGTGATGTGCACGTAAAGTACTTGTAGTCCACTTATAGCCTCTGTCTGGCAATGC
CACAGCCCTGTCAGCATGAATTTGTAATGTCTTGAGCTCTATTATGAATGTGAAGCCTTCCCCTTATCCTCCCTG
TAACCTGATCCATTCTAATTATGTAGCTCTTTGTGAGGGAGTGTTCCCTATCCAATCAATCTTGCATGTAACGC
AAGTTCCCAGTTGGAGCTCCAGCCTGACATCAAAAAGGCAGTTACCATTAAACCATCTCCCTGGTGCTTATGCT
CTTAATTGCCACCTCTAACAGCACCAATCAAAATCTCTCCACTTTCAGCTGTCTTTTGGAGGACGTACGTAATA
AGGTTTTAATTTAGTAAACCAATCCTATGCATGGTTTCAGCACTAGCCAAACCTCACCAACTCCTAGTTCTAGAA
AAACAGGCACCTTGGCAGCCTTGTGATGTCATACAGAGAAGTCACAGGGCAGTACCTGAGGGTCTGTAGGTTGCAC
ACTTTGGTACCAGATAACTTTTTTTTTTCTTTATAAGAAAGCCTGAGTACTCCACACTGCACAATAACTCCTCCC
AGGGTTTTTAACCTTTGTTTTTATTTTCAAACCAGGTCCAATGAGCTTTCTGAACAGCTGGTGTAGCTACAGAGAAA
CCAGCTTCCCTTCAGAGAGCAGTGCTTTTGGCGGGGAGGAGGAAATCCCTTCATACTTGAACGTTTTCTAATTGCT
TATTTATTGTATTCTGGGGTATGGCGTAAGTACAGAGAAGCCATCACCTCAGATGGCAGCTTTTAAAGATTTTT
TTTTTTCTCTCAACACCATGATTCTTTAACAACATGTTTCCAGCATTCCCAGGTAGGCCAAGGTGTCCTACAG
AAAAACCTTGGGTTAGACCTACAGGGGCTCTGGCTGGTGTAAACAGAAGGGAGGGCAGAGCTGGTGCGGCTGGCC
ATGGAGAAAGCTGACTTGGCTGGTGTGGTACAGAGAAGCCAGCTTGTTTACATGCTTATTCCATGACTGCTTGCC
CTAAGCAGAAAGTGCCCTTTCAGGATCTATTTTTGGAGGTTTATTACGTATGTCTGGTTCTCAATTCCAACAGTTT
AATGAAGATCTAAATAAAATGCTAGGTTCTACCTTAAAAAAAAAAAAAAAAAAAA

869/6881
FIGURE 809

MNGQLNGFHEAFIEEGTFLFTSESVGEGHPDKICDQISDAVLDAHLQQDPDAKVACETVAKTGMILLAGEITSRA
AVDYQKVVREAVKHIGYDDSSKGFYKTCNVLVALEQQSPDIAQGVHLDRNEEDIGAGDQGLMFGYATDETEECM
PLTIVLAHKLNAKLAELRRNGTLPWLRPDSKTQVTVQYMQDRGAVLPPIRVHTIVISVQHDEEVCLDEMARDALKEK
VIKAVVPAKYLDDETIYHLQPSGRFVIGGPQGDAGLTGRKIIVDITYGGWGAHGGGAFSGKDYTKVDRSAAYAARW
VAKSLVKGGLCRRVLVQVSYAIGVSHPLSISIFHYGTSQKSERELLEIVKKNFDLRPGVIVRDLDLKKPIYQRTA
AYGHFGRDSFPWEVPKKLKY

[illegible]

871/6881

FIGURE 811

MKHIPVLEDGPWKIVCVKELNGLKKLKRKGKEPARRANGYKTFRLDLEAPEPRAVATNGLRDRTHRLQPVPVPVP
VPVPVAPAVPPRGDTAGERRGGSRAPEVSDARKRCFALGAVGPGLP T P P P P P P P P A P Q S Q A P G G P E A Q P F R E P G P
RPRILLCAPFARPAPSAPPAPPAPPESTVRPAPPTRPGESSYSSISHVIYNNHQDSSASPRKRPGEATAASSEIK
ALQQTRRLLANARERTRVHTISA AFEALRKQVPCYSYGQKLSKLAILRIACNYILSLARLADLDYSADHSNLSFS
ECVQRCTR TLQAEGRAKKRKE

872/6881
FIGURE 812

GGCACGAGGGATGGCGGTTGTATCTGCTGTTTCGCTGGCTGGGCCTCCGCAGCAGGCTTGGCCAGCCGCTGACGGG
TCGGCGGGCGGGTTTGTGTGAACAGGCACGCAGCTGCAGATTTTATTCTGGTAGTGCAACCCCTCTCAAAGGTTGA
AGGAAGTGTATGAACAGGGATTGAAGAAGTAGTAATTCCAAAAAGAAAACCTGGGATAAAGTAGCCGTTCTTCA
GGCACTTGATCCACAGTAAACAGGGATACACAGCTGTGCCTTATGTGTTTCAAGATGATCCTTACCTTATGCC
AGCATCATCTTTGGAATCTCGTTCATTTTTACTGGCAAAGAAATCCGGGGAGAATGTGGCCAAGTTTATTATTAA
TTCATACCCCAAATATTTTTCAGAAGGACATAGCTGAACCTCATATACCGTGTTTAAATGCCTGAGTACTTTGAACC
TCAGATCAAAGACATAAGTGAAGCCGCCCTGAAGGAACGAATTGAGCTCAGAAAAGTCAAAGCCTCTGTGGACAT
GTTTGATCAGCTTTTGCAAGCAGGAACCACTGTGTCTCTTGAACAACAAATAGTCTCTTGGATTTATTGTGTTA
CTATGGTGACCAGGAGCCCTCAACTGATTACCATTTTCAACAACTGGACAGTCAGAAGCATTGGAAGAGGAAAA
TGATGAGACATCTAGGAGGAAAGCTGGTCATCAGTTTGGAGTTACATGGCGAGCAAAAAACAACGCTGAGAGAAT
CTTTTCTCTAATGCCAGAGAAAAATGAACATTCCATTGACAAATGATCCGAGGAATGGTGAAGCACCAGCTTA
TGAGCAGGCATTAACTTGTACACTGAGTTACTAAACAACAGACTCCATGCTGATGTATACACATTTAATGCATT
GATTGAAGCAACAGTATGTGCGATAAATGAGAAATTTGAGGAAAAATGGAGTAAATACTGGAGCTGCTAAGACA
CATGGTTGCACAGAAGGTGAAACCAAATCTTCAGACTTTTAAATACCATTCTGAAATGTCTCCGAAGATTTTATGT
GTTTGCAAGATCGCCAGCCCTTACAGGTTTACGTGAAATGAAAGCCATTGGAATAGAACCCTCGCTTGCAACATA
TCACCATATTATTCGCCCTGTTTGATCAACCTGGAGACCCCTTAAAGAGATCATCCTTCATCATTATGATATAAT
GAATGAATTAATGGGAAAGAGATTTTCTCAAAGGACCCGGATGATGATAAGTTTTTTCAGTCAGCCATGAGCAT
ATGCTCATCTCTCAGAGATCTAGAACTTGCTACCAAGTACATGGCCTTTTAAAAACCGGAGACAACCTGGAAAT
CATTGGACCTGATCAACATCGTAATTTCTATTATTCCAAGTTCTTCGATTTGATTGTCTAATGGAACAAATTGA
TGTTACCTTGAAGTGGTATGAGGACCTGATACCTTCAGCCTACTTTCCCACTCCCAAACAATGATACATCTTCT
CCAAGCATTGGATGTGGCCAATCGGCTAGAAGTGATTCTAAAATTTGGAAAGATAGTAAAGAATATGGTTCATAC
TTTCCGCGAGTGACCTGAGAGAAGAGATCCTGATGCTCATGGCAAGGGACAAGCACCCACCAGAGCTTCAGGTGGC
ATTTGCTGACTGTGCTGCTGATATCAAATCTGCGTATGAAAGCCAACCCATCAGACAGACTGCTCAGGATTGGCC
AGCCACCTCTCTCAACTGTATAGCTATCCTCTTTTAAAGGGCTGGGAGAACTCAGGAAGCCTGGAAAATGTTGGG
GCTTTTCAGGAAGCATAATAAGATTCTTAGAAGTGAGTTGCTGAATGAGCTTATGGACAGTGCAAAGTGTCTAA
CAGCCCTTCCCAGGCCATTGAAGTAGTAGAGCTGGCAAGTGCCCTTCAGCTTACCTATTTGTGAGGGCCTCACCCA
GAGAGTAATGAGTGATTTTGCAATCAACCAGGAACAAAAGGAAGCCCTAAGTAATCTAACTGCATTGACCAGTGA
CAGTGATACTGACAGCAGCAGTGACAGCGACAGTGACACCAGTGAAGGCAAATGAAAGTGGAGATTGAGGAGCAG
CAATGGTCTCACCATAGCTGCTGGAATCACACCTGAGAACTGAGATATACCAATATTTAACATTGTTACAAAGAA
GAAAAGATACAGATTTGGTGAATTTGTTACTGTGAGGTACAGTCAGTACACAGCTGACTTATGTAGATTTAAGCT
GCTAATATGCTACTTAACCATCTATTAATGCACCATTAAGGCTTAGCATTTAAGTAGCAACATTGCGGTTTTCA
GACACATGGTGAGGTCCATGGCTCTTGTATCAGGATAAGCCTGCACACCTAGAGTGTGCGGTGAGCTGACCTCAC
GATGCTGTCTCGTGCGATTGCCCTCTCTGCTGCTGGACTTCTGCCTTTGTTGGCCTGATGTGCTGCTGTGATG
CTGGTCTTCATCTTAGGTGTTTATGCAGTTCTAACACAGTTGGGGTTGGGTCAATAGTTTCCCAATTTGAGGAT
ATTTGATGTGAGAAATAACGCATCTTAGGAATGACTAAACAAGATAATGGCAGTTTAGGCTGCACAACCTGGTAA
AATGACTGTAGATAAATGTTGTAATTAGTGTACAGTTTGTATTTTTGTTAATATAGCCGCTGCCATAGTTTTCT
AACTTGAACAGCCAAAAA

873/6881
FIGURE 813

MVAQKVKPNLQTFNTILKCLRRFHV FARSPALQVLREMKAIGIEPSLATYHHIIRLFDQPGDPLKRSSFIIYDIM
NELMGKRFSPKDPDDDKFFQSAMSSICSSLRDLELAYQVHGLLKTGDNWKFIGPDQHRNFYYSKFFDLICLMEQID
VTLKWYEDLIPSAYFPHSQTMIHLLQALDVANRLEVIPKIWKDSKEYGHTFRSDLREEILMLMARDKHPPPELQVA
FADCAADIKSAYESQPIRQTAQDWPATSLNCIAILFLRAGRTQEAWKMLGLFRKHNKIPRSELLNELMDSAKVSN
SPSQAIIEVVELASAFSLPICEGLTQRVMSDFAINQEQKEALSNLTALTSDDTDSSSDSDSDTSECK

874/6881

FIGURE 814

ACGCGGGCACGCACACACGGAAGCACGCCTCCACTTAACTCGCGCCGCCGCGGCAGCTCGAGTCCACCAGCAGCG
CCGTCCGCTTGACCGAGATGCTGCGGGCCTGTCAGTTATCGGGTGTGACCGCCGCCGCCAGAGTTGTCTCTGTG
GGAAGTTTGTCTCCGTCCATTGCGACCATGCCGAGATACTCTACTTCAGGCAGCTCTGGGTTGACTACTGGCA
AAATTGCTGGAGCTGGCCTTTTGTGTTGTTGGTGAGGTATTGGTGGCACTATCCTATATGCCAAATGGGATTCCC
ATTTCCGGGAAAGTGTAGAGAAAACCATACCTTACTCAGACAACTCTTCGAGATGGTCTTGGTCTGCAGCTT
ATAATGTTCCATTGCCAAAGAAATCGATTTCAGTCGGGTCCACTAAAAATCTCTAGTGTATCAGAAGTAATGAAAG
AATCTAAACAGTCTGCCTCACAACTCCAAAAACAAAAGGGAGATACTCCAGCTTCAGCAACAGCACCTACAGAAG
CGGCTCAAATTATTTCTGCAGCAGGTGATACCTGTGCGTCCCAGCCCCTGCAGTTCAGCCTGAGGAATCTTTAA
AAACTGATCACCTGAAATTGGTGAAGGAAAACCCACACCTGCACCTTCAGAAGAAGCATCCTCATCTTCTATAA
GGGAGCGACCACTGAAGAAGTTGCAGCTCGCCTTGCAACAGGAAAAACAAGAACAAGTTAAAATTGAGTCTC
TAGCCAAGAGCTTAGAAGATGCTCTGAGGCAAACCTGCAAGTGTCACTCTGCAGGCTATTGCAGCTCAGAATGCTG
CGGTCCAGGCTGTCAATGCACACTCCAACATATTGAAAGCCGCCATGGACAATTCTGAGATTGCAGGCGAGAAGA
AATCTGCTCAGTGGCGCACAGTGGAGGGTGCATTGAAGGAACGCAGAAAGGCAGTAGATGAAGCTGCCGATGCC
TTCTCAAAGCCAAAGAAGAGTTAGAGAAGATGAAAAGTGTGATTGAAAATGCAAAGAAAAAGAGGTTGCTGGGG
CCAAGCCTCATATAACTGCTGCAGAGGGTAAACTTCACAACATGATAGTTGATCTGGATAATGTGGTCAAAGAGG
TCCAAGCAGCTCAGTCTGAGGCTAAGGTTGTATCTCAGTATCATGAGCTGGTGGTCCAAGCTCGGGATGACTTTA
AACGAGAGCTGGACAGTATTACTCCAGAAGTCCTTCTCGGATGGAAGGAATGAGTGTTCAGACTTAGCTGACA
AGCTCTCTACTGATGATCTGAACCTCCTCATTGCTCATGCACATCGTCGTATTGATCAGCTGAACAGAGAGCTGG
CAGAACAGAAGGCCACCGAAAAGCAGCACATCACGTTAGCCTTGGAGAAAACAAAAGCTGGAAGAAAAGCGGGCAT
TTGACTCTGCAGTAGCAAAAGCATTAGAACATCACAGAAGTGAAATACAGGCTGAACAGGACAGAAAGATAGAAG
AAGTCAGAGATGCCATGGAAAATGAAATGAGAACCCAGCTTCGCCGACAGGCAGCTGCCCACACTGATCACTTGC
GAGATGTCCTTAGGGTACAAGAACAGGAATTGAAGTCTGAATTTGAGCAGAACCTGTCTGAGAACTCTCTGAAC
AAGAATTACAATTCGTCGTCTCAGTCAAGAGCAAGTTGACAACCTTTACTCTGGATATAAATACTGCCTATGCCA
GACTCAGAGGAATCGAACAGGCTGTTTCAGAGCCATGCAGTTGCTGAAGAGGAAGCCAGAAAAGCCCACTCT
GGCTTTTCAGTGGAGGCATTAAAGTACAGCATGAAGACCTCATCTGCAGAAACACCTACTATCCCCTGGGTAGTG
CAGTTGAGGCCATCAAAGCCAAGTGTCTGATAATGAATTCACCCAAGCTTTAACCGCAGCTATCCCTCCAGAGT
CCCTGACCCGTGGGGTGTACAGTGAAGAGACCCTTAGAGCCGTTTCTATGCTGTTCAAAAAGTGGCCGAAGGG
TAGCAATGATTGATGAAACCAGAAATAGCTTGTACAGTACTTCTCTCCTACCTACAGTCCCTGCTCCTATTCC
CACCTCAGCAACTGAAGCCGCCCCCAGAGCTCTGCCCTGAGGATATAAACACATTTAAATTACTGTCATATGCTT
CCTATTGCATTGAGCATGGTGATCTGGAGCTAGCAGCAAAGTTTGTCAATCAGCTGAAGGGGGAATCCAGACGAG
TGGCACAGGACTGGCTGAAGGAAGCCGAATGACCCTAGAAACGAAACAGATAGTGGAAATCCTGACAGCATATG
CCAGCGCCGTAGGAATAGGAACCACTCAGGTGCAGCCAGAGTGGAGGTTAGGAAGATTTTCATAAAGTCATATTT
CATGTCAAAGGAAATCAGCAGTGATAGATGAAGGGTTCGCAGCGAGAGTCCCGGACTTGTCTAGAAATGAGCAGG
TTTACAAGTACTGTTCTAAATGTTAACACCTGTTGCATTTATATCTTTCCATTTGCTATCATGTGAGTGAACGC
CAGGAGTGCTTTCTTTGCAACTTGTGTAACATTTCTGTTTTTTCAGGTTTTACTGATGAGGCTTGTGAGGCCAA
TCAAAATAATGTTTGTGATCTCTACTACTGTTGATTTTGCCCTCGGAGCAAACCTGAATAAAGCAACAAGATG

875/6881

FIGURE 815

MLRACQLSGVTAAAQSCLCGKFVLRPLRPCRRYSTSGSSGLTTGKIAGAGLLFVGGGIGGTILYAKWDSHFRESV
EKTIPYSDKLFEMVLGPAAYNVPLPKKSIQSGPLKISSVSEVMKESKQSASQLQKQKGDTPASATAPTEAAQIIS
AAGDTLSVPAPAVQPEESLKTDPHPEIGEGKPTPALSEEASSSSIRERPPEEVAARLAQQEKQEQVKIESLAKSLE
DALRQTASVTLQAIQAQNAAVQAVNAHSNLIKAAAMDNSEIAGEKKSAQWRTVEGALKERRKAVDEAADALLKAKE
ELEKMKSVIENAKKKEVAGAKPHITAAEGKLHNMIVDLDNVVKKVQAAQSEAKVVSQYHELVVQARDDFKRELD
ITPEVLPGWKGMSSVDLADKLSTDDLNSLIAHAHRRIDQLNRELAEQKATEKHITLAEKQKLEEKRAFD
KALEHHRSEIQAEQDRKIEEVRDAMENEMRTQLRRQAAHTDHLRDVLRVQEQELKSEFEQNLSEKLSEQELQFR
RLSQEQVDNFTLDINTAYARLRGIEQAVQSHAVAEAEARKAHQLWLSVEALKYSMTSSAETPTIPLGSAVEAIK
ANCSDNFTQALTAIIPPELSTRGVYSEETLRARFYAVQKLARRVAMIDETRNSLYQYFLSYLSLLFPQQLK
PPPELCPEDINTFKLLSYASYCIEHGDLELAAKFVNQLKGESRRVAQDWLKEARMILETKQIVEILTAYASAVGI
GTTQVQPE

876/6881
FIGURE 816

TAGTGCCGCCGCTCCCTACTCTGGGGGTTGGGACTACCTCCTTTTCCGCGGGCCCCGCCAGGCGGCTGCCCCGTG
ACCTGCCTGGGCGCGGGGAACTGAAAGCCGGAAGGGGCAAGACGGGTTTCAGTTCGTTCATGGGGCTGTTTGAAAG
ACCCAGGAGAAGCCGCCCAAAGAACTGGTCAATGAGTGGTCATTGAAGATAAGAAAGGAAATGAGAGTTGTTGAC
AGGCAAATAAGGGATATCCAAAGAGAAGAAGAAAAAGTGAACCGATCTGTGAAAGATGCTGCCAAGAAGGGCCAG
AAGGATGTCTGCATAGTTCTGGCCAAGGAGATGATCAGGTCAAGGAAGGCTGTGAGCAAGCTGTATGCATCCAAA
GCACACATGAAGTCAAGTGCATGGGGATGAAGAACCAGCTCGCGGTCTTGCGAGTGGCTGGTTCCCTGCAGAAG
AGCACAGAAGTGATGAAGGCCATGCAAAGTCTTTGTGAAGATTCCAGAGATTTCAGGCCACCATGAGGGAGTTGTCC
AAAGAAATGATGAAGGCTGGGATCATAGAGGAGATGTTAGAGGACACTTTTGAAAGCATGGACGATCAGGAAGAA
ATGGAGGAAGAAGCAGAAATGGAAATTGACAGAATTCTCTTTGAAATTACAGCAGGGGCCTTGGGCAAGCACCC
AGTAAAGTGACTGATGCCCTTCAGAGCCAGAACCTCCAGGAGCGATGGCTGCCTCAGAGGATGAGGAGGAGGAG
GAAGAGGCTCTGGAGGCCATGCACTCCCGGCTGGCCACACTCCGCGAGTAGGGGCTGCCTACCCCGCTGGGTGTG
CACACACTCCTCTCAAGAGCTGCCATTTTATGTGTCTCTTGCACTACACCTCTGTTGTGAGGACTACCATTTTGG
AGAAGGTTCTGTTGTCTCTTTTCATTCTCTGCCAGGTTTTGGGATCGCAAAGGATTGTTCTTATAAAAGTGG
CATAAATAAATGCATCATTTTTAGGAGTATAGACAGATATATCTTATTGTGGGAGGGGAAAGAAATCCATCTGC
TCATGAAGCACITCTGAAAATATAGGTGATTGCCTGAATGTGCAAGACTCTACTTTTGTCTATAAAACACTATAT
AAATGAATTTTAAATAAATTTTGTCTTAGCACTTGGCCCCATTGTAGATTGCCCTGTGCAGTAAACTTTCAAGGT
GTCGGCTGCCCCAGATTGCTTCATTGCTGGGTGTGGAAAGAGTTGCTATGGCCAGGCATATGGGATTGTGAAGC
TCAGCAGAAGTGACTTCTGCTCTGTGGTTGCTGCTCCCCGGCTTTACAGACATGGTATGGCAGCCATTCTTTTA
TCTATTTAACCAAGAGGATGCTGGGGAATTGTGCTGCTTGTCTGTTGGCTGGTGGCTGCATTATGTCTGGGGT
GTGCATGTGGGTCTATTTAGAGCTTCTGTCCCTTCCTTCCCATTGCAAGTTGCACCCAGATGAGACAGCTGTAGT
ACTAGGTCTCTTTACCTCTCATTGCTGTCCCTGCTTCGAGCTGGTTGTCTTGTGCGTGGGACATGGGCCTTCC
TATCTGTGTTTTCTCAAAGTCAGGAGCTGACCAGGAGCACACTAAGGTGTGGTCATGCATCATAACCAACATTCA
CTCATCTGGGACATTCTTAAGATACATTATAAATCATTTCAGCAGTAGTACTTTGTATGTGTTGAGAGTTTACA
GAGCTCTTTGACATACGCGATCTTAGTCTTTACAAATAAGGAAAACAGCTCAGTTTGGGAAGTATCAGAGATGGG
ATTCAAACCCAGATCCTCTGGTCCAAGTTGTATGTGCACTGAACATAATCAGGCAGGAAAAAAGCCAGCCACTGT
CTCAGAGATTGTTTTTGTATATTGTAGCAAAATCCTGAAACAATGGGGTCCCTCCAGTCTCATCATACAAATG
GCAATCTTGGCTGGGTGCGGTGGTTTCATGCCTATAATCCCAGTGCTTTTACAAGGCTGAGGCAGGAGGCTCTCTTG
AGAATAGGAGTTCAAGACCAGCCTGGGCAACATAGCAAGATCCTGTCTCTCCAAAAAAGGCTGCTCTCTG
AAAATTTCATTTTTGTAGTCCAGAGGACCTCCTATTACTCTTGATTTTCATCTTCAGAGTGTAGTTAAAAAATTAT
TTTAAATAAATTATTTTTTAAATCAGTTGTAGGTTTACAGCAAAAGTGGACAAAAAGAAATTTCTCATATATCCC
CTGCCCTCACACATGCATAGCCTCCCACCACTATCAGTATCCACACCAGAGTGGTACATTTGTTACAATCAATA
AACCTCCATTGACACATCATTATCACCCAAAGTCCATAGTTTACATGAAGATTCACTCTGGTGTGTACATTGTA
TGGGCTTAGACAAATGTATGATGATATCTACAATTATAGAATCATACAGAATAGTTTCACTGCCCTAAAACTTCT
CTATGCTTCACCTGTTTCATCCCTTTCTTCCCTAATCCCCCTGGCAACCACTTTAAAAAAGGCTTATAGGTTTCAAGG
GTACATGTGCAGGTAACTCGTGACAAGGGGGTTGTTATACAGATTATTTAGTGACCCAGGTACTAAGCCTAGT
ACCCAATAGTTACTTTTCTGGTCCCTGTCCCTTTTCCACCCCTCAGGTAGGCCCCAGTATGTTATTCCT
TTGTGTCCATGTTATTTCACTCCCACTTGTGAGAACATGGAATATTGGTTTCTGTTTCTATGTTAGTTTGTTA
AGGATAATGGCCTCCAGCCCCATCCATGTTCCCTGCAAGGACATGATCTTTCCTTGGCAACCACTTTTACTGTC
GCCATAGTTCTTCTTTCTAGAATGTATATTGGAATCATATAGTATGTAGCCTTTTCAGACTGGCTTCTTCA
CTTAATAATATGCAATTAAGGTTCCCTCCATGTCAATTTCATGGCTTAATAGTGCATTTATTTTAGCACTGAATAA
TACTCCATTGTCTAGATGAATAGTTTATCCATTACCTATTGAAAGACTTCTTGGTGGTTTCCAAGTTTGGCAA
TTATGAATAAAGCTGTTGTAAACATCTTTGTGCAGGTTTTCTATGGGCATGTTTTTAATTCATTTGAATAAATA
CCAAGAGCTTCAGTGCTGGATCATA

877/6881
FIGURE 817

CAGGACACAGCATGGACATGAGGGTCCCCGCTCAGCTCCTGGGGCTCCTGCTGCTCTGGCTCCCAGGTGCCAAAT
GTGACATCCAGATGACCCAGTCTCCTTCCACCCTGTCTGCATCTGTAGGAGACAGAGTCACCATCACTTGCCGGG
CCAGTCAGAGTATTAGTAGCTGGTTGGCCTGGTATCAGCAGAAACCAGGGAAAGCCCCCTAAGCTCCTGATCTATA
AGGCGTCTAGTTTAGAAAGTGGGGTCCCATCAAGGTTAGCGGCAGTGGATCTGGGACAGAATCACTCTCACCA
TCAGCAGCCTGCAGCCTGATGATTTTGCAACTTATTACTGCCAACAGTATAATAGTTATTCTCCACATGGACGT
TCGGCCAAGGGACCAAGGTGGAAATCAAACGAACGTGGCTGCACCATCTGTCTTCATCTTCCCGCCATCTGATG
AGCAGTTGAAATCTGGAAGTGCCTCTGTTGTGTGCCTGCTGAATAACTTCTATCCCAGAGAGGCCAAAGTACAGT
GGAAGGTGGATAACGCCCCCTCCAATCGGGTAACTCCCAGGAGAGTGTACAGAGCAGGACAGCAAGGACAGCACCT
ACAGCCTCAGCAGCACCCCTGACGCTGAGCAAAGCAGACTACGAGAAACACAAAGTCTACGCCTGCGAAGTCACCC
ATCAGGGCCTGAGCTCGCCCGTCACAAAGAGCTTCAACAGGGGAGAGTGTTAGAGGGAGAAGTGCCCCCACCTGC
TCCTCAGTTCCAGCCTGACCCCTCCCATCCTTTGGCCTCTGACCCTTTTCCACAGGGGACCTACCC

878/6881
FIGURE 818

GCGGCCGTTACGGCGCTCAGGCGTCTCGACGCGCGGATTTAAACCAGCTCAGGAGACGCCAAGGAAAGATGGG
ACCTCCCGGCCAGCACTGCCAGCCACAATGAATAACTCTTCTTCAGAGACGCGAGGACACCCCCACAGTGCCTC
CTCTCCTTCAGAGCGTGTGTTCCCGATGCCCCTGCCAGGAAGGCGCTCTCAATATTCCTGGCACCCAGTCTCT
CGAAGACTTTCCTCAGAATGACGATGAGAAGGAGCGGCTGCAGCGGAGGCGCTCGAGGGTCTTTGATCTCGAGTT
CAGCACTGACTCAGCTCGCTTATTGGCCTCCCCCTCCAGCAGGAGTATTGACATTTAGCTACTATCCCCAAGTT
TACAAACACGCAGATTACGGAACATTACTCCACCTGTATCAAACCTGTCCACTGAAAATAAAATCACTACCAAGAA
TGCTTTTGGTTTGCACTTGATTGATTTTATGTCTAGAGATTCTTAAACAGAAAGACACCGAACCAACCACTTTAA
AGTGGCTGCGGGTACTCTGGATGCCAGCACCAAGATCTATGCTGTGCGCGTGGATGCCGTCCATGCCGATGTATA
CAGAGTCTTTGGGGGCTGGGCAAAGATGCACCGTCTTTGGAAGAAGTAGAAGGCCATGTTGCTGATGGAAGTGC
TACTGAAATGGGAACAACCAAAAAGGCTGTAAAGCCAAAGAAGCACTTACACAGAACTATTGAGCAGAACAT
AAACAACCTCAATGTCTCCGAAGCAGATCGGAAGTGTGAGATTGATCCCATGTTTCAGAAGACAGCAGCTCATT
TGATGAGTGCAGCACAGCAGGGGTGTTTCTGTCCACTCTCCACTGCCAGGACTACAGAAGTGAAGTGTGTTTCC
CTCTGATGTCCAGACTCTCTCCACGGGAGAACCTCTCGAGTTGCCAGAGTTAGGTTGTGTAGAAATGACAGATT
AAAAGCGCCCTTGCAGCAGTGTGCAGAAGATCGCCAGATCTGCCCTTCCCTGGCCGGGTTCAGTTTACACAGTG
GGACAGTGAAACACATAATGAGTCTGTGTGCGGCCCTGGTAGACAAGTTTAAAGAAGATGACCAGGTATTTGACAT
CAATGCTGAAGTTGACGAGAGTGAAGTGTGGAGACTTCCCGATGGGTCCCTGGGGGATGACTTTGATGCCAACGA
TGAACCTGACCACACCGCAGTTGGGGATCATGAAGAGTTCAGGAGCTGGAAGGAGCCCTGCCAGGTTGAGAGCTG
CCAGGAAGAAATGATTTCCCTTGGGGATGGAGACATCAGGACCATGTGCCCTTCTGTCTATGAAACCTGGAGA
ATATTTCTATTTTCACTCTCGGACCATGTCTGATGTGGGTGGCCCGGATCACTGGCGCTTTAGGCTCGACGCAA
ACAAGATGCTCCTTCCCAATCAGAAAACAAAAAGAAGAGTACAAAAAAGATTTGAAATTGACTTTGAAGATGA
TATTGACTTTGATGTATATTTTAGAAAAACAAAGGCTGCTACTATTCTGACCAAGTCCACTTTGGAGAACCAGAA
TTGGAGAGCTACCACCTTCTACAGATTTCAACTACAATGTTGACACTCTGGTCCAGCTTACCTCAAACCAGG
CACCAGGTTACTTAAGATGGCCCAGGGCCATAGGGTAGAGACTGAGCATTATGAAGAAATTGAAGACTATGATTA
CAACAACCTTAACGACACCTCCAACCTTTTGCCCTGGATTACAGGCTGCTGACAGTGATGATGAAGATTGGATGA
CTTATTTGTGGGACCTGTTGGGAACTCTGACCTCTCACCTTATCCTTGCCATCCACCTAAGACAGCACAACAGAA
TGGTGACACTCCAGAAGCCCAAGGATTAGACATCACAACATATGGGGAGTCAAACCTGGTAGCTGAGCCTCAGAA
GGTAAATAAAATTGAAATTCACTATGCCAAGACTGCCAAAAGATGGACATGAAGAACTGAAGCAGAGCATGTG
GAGTCTGCTGACAGCGCTCTCCGGAAAGGAGGCAGATGCAGAGGCAAACCACAGGGAAGCTGGAAAAGAAGCGGC
CCTGGCAGAAAGTGGCTGACGAGAAGATGCTTAGCGGGCTCACGAAGGACCTGCAGAGGAGCCTGCCCCCTGTCTAT
GGCTCAGAACCTCTCCATACCTCTGGCTTTTGCTGTCTCCTACATTTAGCCAATGAAAAGAACTTAAACTGGA
AGGAACAGAGGACCTCTCTGATGTTCTTGTGAGGCAAGGAGATTGAGTTCACTATGGAGAAGTCAGCAGCAGGAG
GCCATCCCTTACTCAGTTGCCGGGACATCCCCAGTCTCGGGGAAGAAGATGCCATGGGCTTATACCCAGGCTG
TAGCCAACACCAACGTGCCTGTTTGTGTTGTGCTCTTCTCTCTCCATCATAGTCTGGGTGCCAGCGCCCTG
AAGCTCCGTGCTCAACTGATTAACTTTACTGCCCTATGGTGACCATCTAGGAGAGGGGAGGGCAGAGGGGTGA
GGGTACTATTCTGGATTGAGAAAACCTATATCCATTCTTTATATCAATGTATAGTTTTAGTCTCCTAAATTGATC
TGTTATTTTCCAACTATTCTCTTGTAGAAAATTTCCAGTGGGCACCTAATGGTGCCCTTGAAGAAGCTTCTTAA
TCCATGTACATAAAATACATCATATGTACACTTATAAATGTATATAGAATGCTCAAAAATAAAATTCTTAATAAT
AGAACTGGCAAAATA

879/6881
FIGURE 819

MGPPGPALPATMNNSSSETRGHPSASSPSERVFPMLPRKAPLNIPGTPVLEDFPQNDDEKERLQRRRSRVFDL
QFSTDSPRLLASPSRSIDISATIPKFTNTQITEHYSTCIKLSTENKITTNAFGLHLIDFMSEILKQKDTEPTN
FKVAAGTLDASTKIYAVRVDAVHADVYRVLGGLGKDAPSLEEVEGHVADGSATEMGTTKAVKPKKKHLHRTIEQ
NINNLNVSEADRKCEIDPMFQKTAASFDECSTAGVFLSTLHCQDYRSELLFPSDVQTLSTGEPLLELPELGCVEMT
DLKAPLQQCAEDRQICPSLAGFQFTQWDSETHNESVSALVDKFKKNDQVFDINAEVDESDCGDFPDGSLGDDFDA
NDEPDHTAVGDHEEFRSWKEPCQVQSCQEEMISLGDGDIRTMCPLLSMKPGEYSYFSPRTMSMWAGPDHWRFRPR
RKQDAPSQSENKKKSTKKDFEIDFEDDIDFDVYFRKTKAATILTKSTLENQNWRAATLPTDFNYNVDTLVQLHLK
PGTRLLKMAQGHRVETEHYEEIEDYDYNPNPNDTSNFCPLQAADSDDDLDDLFVGPVGNSDLSPYPCHPPKTAQ
QNGDTPEAQGLDITTYGESNLVAEPQKVNKIEIHYAKTAKKMDMKKLQSMWSLLTALSGKEADAEANHREAGKE
AALAEVADEKMLSGLTQDLQSLPVPMAQNLSIPLAFACLLHLANEKNLKLKLEGTEDLSVDLVLRQGD

880/6881
FIGURE 820A

TTCCAAGAAGAAGGATTTGCACCCCTCGGGATATTGATGCATTTTGGCTGCAGCGGCAGCTCAGTCGTTTCTATGA
TGATGCCATCGTGTGCGCAGAAGAAGGCAGATGAAGTATTGGAGATTTTGAAGACGGCCAGTGATGATCGGGAATG
TGAAATCAGCTGGTTCTGCTGCTTGGTTTCAACACCTTTGATTTTCATTAAAGTGTTCGGGCAGCACAGGATGAT
GATTTTATACTGTACCTTGCTGGCCAGTGCACAAAGTGAAGCTGAAAAGGAAAGGATTATGGGAAAGATGGAAGC
TGACCCAGAGCTATCCAAGTTCCTCTACCAGCTTCATGAAACCGAGAAGGAGGATCTGATCCGAGAGGAAAGGTC
CCGGAGAGAGCGAGTGCCTCAGTCTCGAATGGACACAGATCTGGAACCATGGATCTCGACCAGGGTGGAGAGGC
ACTGGCTCCACGGCAGGTTCTGGACTTGGAGGACCTGGTTTTTACCCTAAGGGAGCCACTTTATGGCCAATAAACG
CTGTCAGCTTCTGATGGATCCTTCCGTGCGCCAGCGTAAGGGCTATGAAGAGGTGCATGTGCCTGCTCTGAAGCC
CAAGCCCTTTGGCTCAGAAGAACAACCTGCTTCCAGTGGAAAAGCTGCCAAAGTATGCCAGGCTGGGTTTGAGGG
CTTCAAAACACTGAATCGGATCCAGAGTAAGCTCTACCGTGCTGCCCTTGAGACGGATGAGAATCTGCTGCTGTG
TGCTCCTACTGGTGTGGGAAGACCAACGTGGCCCTGATGTGCATGCTCCGAGAGATTGGGAAACACATAAACAT
GGACGGCACCATCAATGTGGATGACTTCAAGATTATCTACATTGCCCCCATGCGCTCCTTGGTGCAGGAGATGGT
GGGCAGCTTTGGAAAGCGCCTGGCCACTTATGGCATCACTGTTGCTGAACTGACTGGGGACCACAGCTGTGCAA
AGAAGAGATCAGTGGCACTCAGATCATCGTCTGCACCCCGAGAAGTGGGACATCATACCCGCAAGGGTGGTGA
GCGACCTACACCCAGCTGGTGGGGCTCATCTTCTGGATGAGATTCTTCTCCACGATGACAGAGGTCTGT
CTTAGAAGCTTTAGTGGCCAGGGCCATCCGAAACATTGAGATGACCCAAGAGGATGTCCGACTCATTGGTCTCAG
TGCCACCCCTACCCAACATATGAAGATGTAGCCACCTTTCTACGTGTTGACCCTGCCAAGGGTCTCTTTTACTTTGA
CAACAGCTTCCGTCCAGTGCCTCTGGAACAGACATATGTGGGTATCACAGAGAAAAAGCTATCAAGCGTTTCCA
GATCATGAATGAATCGTCTATGAAAAATCATGGAACATGCTGGAAAAATCAGGTGCTGGTGTGTTGTCCACTC
CCGGAAGGAGACTGGAAAGACAGCCAGGGCCATCCGGGACATGTGCCTAGAAAAGGACACTCTGGGTCTGTTTCT
GAGGGAGGGCTCAGCCTCCACAGAAGTCTGCGAACAGAAGCTGAGCAGTGCAAGAACCTAGAGCTGAAGGATCT
TCTGCCCTTATGGCTTTGCTATTATCACGCAGGCATGACCAGGGTTGACCGAACACTCGTGAGGATCTTTTTGC
TGATAAACATATTAGGTTTTAGTTTCCACAGCAACTCTAGCTTGGGGTGTGAATCTCCCTGCACATACAGTCAT
CATCAAAGGCACCCAGGTGTACAGTCCAGAGAAGGGGCGTTGGACAGAAGTGGGAGCACTGGACATTCTGCAGAT
GCTGGGACGTGCCGGAAGACCCAGTATGACACCAAGGGTGAAGGCATACTCATCATCTCATGGGGAGCTACA
GTACTACCTGTCCCTCCTCAATCAACAACCTTCTATTGAAAGCCAGATGGTTCAAAGCTTCTGACATGCTCAA
TGCAGAAATCGTGTAGGAAATGTCCAGAAATGCCAAGGATGCGGTGAACTGGCTGGGCTATGCCTACCTCTATAT
CCGAATGCTGCGATCCCCAACCTCTATGGCATCTCTCATGATGACCTCAAGGGAGATCCCCTGCTGGACACGG
CCGACTAGATCTGGTTCATACAGCTGCCCTGATGTGTCGACAAGAACAATCTGGTCAAGTACGACAGAAGACGGG
CAACTTCCAGGTGACAGAACTGGGCCGTATAGCCAGCCACTACTACATCACCATGATACAGTGCAGACTTACAA
CCAGCTGCTGAAGCCCACCTGAGTGAGATTGAGCTTTTTCAGGGTCTTCTCATTGTCTCTGAGTTCAAGAACAT
CACAGTGAGAGAGGAGGAGAAGCTGGAGCTGCAGAAGTTGCTGGAGAGGGTGCCTATCCCTGTAAAGGAGAGCAT
TGAGGAACCCAGTGCTAAGATCAACGTTCTTCTGCAAGCCTTCATCTCACAGCTGAAATTGGAGGGCTTTGCACT
GATGGCTGACATGGTGTATGTACACAGTCGGCTGGCCGGTTGATGCGAGCGATAATTGAAATTGTCTGAAACCG
AGGTTGGGCACAGCTTACAGACAAGACCTGAACCTCTGCAAGATGATCGACAAACGCATGTGGCAGTCCATGTG
TCCTCTGCGCCAGTTCCGGAAACTCCCTGAGGAAGTAGTGAAGAAGATTGAGAAGAAGAATTTCCCTTTGAGCG
TCTGTACGACCTGAATCATAATGAGATTGGGGAGCTTATCCGCATGCCAAAGATGGGGAAGACCATCCACAAATA
TGTCATCTGTTTCCCAAGTTGGAGTTGTCTAGTGCACCTGCAGCCTATCACACGCTCCACCCTGAAGGTGGAGCT
GACCATCACGCCAGACTTCCAGTGGGATGAAAAGGTGCATGGTTTCATCCGAGGCTTTTGGATTCTGGTGGAGGA
TGTGGACAGCGAGGTGATTCTGCACCATGAGTATTTCTCCTCAAGGCCAAGTACGCCCAGGACGAGACCTCAT
TACATTCTTCGTGCCTGTCTTTGAACCGCTGCCCCCTCAGTACTTCATCCGAGTGGTGTCTGACCGCTGGCTCTC
TTGTGAGACCCAGCTGCCTGTCTCCTTCCGGCACCTGATCTTGGCGAGAAGTACCCCCCTCCAACCGAATTTT
GGACCTGCAGCCCTTGGCCGTGTCTGCTCTGAGAACAGTGCCTTTGAGAGTCTTTACCAAGATAAATTTCTTTT
CTTCAATCCCATCCAGACCCAGGTGTTTAAACATGTATACAACAGTGACGACAACGTGTTTGTGGGGGCCCCAC
GGGCAGCGGGAAGACTATTTGTGCAGAGTTTGCCATCCTGCGAATGCTGCTGCAGAGCTCGGAGGGGCGCTGTGT
GTACATACCCCCATGGAGGCCCTGGCAGAGCAGGTATACATGGACTGGTACGAGAAGTTCCAGGACAGGCTCAA
CAAGAAGTGGTACTCCTGACAGGCGAGACCAGCACAGACCTGAAGCTGCTGGGCAAAGGGAACATTATCATCAG
CACCCCTGAGAAGTGGGACATACTTTCCCGGCGATGGAAGCAGCGCAAGAACGTGCAGAACATCAACCTCTTCGT

FIGURE 820B

GGTGGATGAGGTCCACCTTATCGGGGGCGAGAATGGGCCTGTCTTAGAAGTGATCTGCTCCCGAATGCGCTACAT
CTCCTCCCAGATTGAGCGGGCCATTTCGCAATTGTGGCACTCAGCTCTTCGCTCTCCAATGCCAAGGATGTGGCCCCA
CTGGCTGGGCTGCAGTGCCACCTCCACCTTCAACTTCCATCCCAATGTGCGTCCCGTCCCTTGGAGCTGCACAT
CCAGGGCTTCAACATCAGCCATACACAAACCGCCTGCTCTCCATGGCCAAGCCTGTGTACCATGCTATCACCA
GCACTCGCCCCAAGAAGCCTGTCTATTGTCTTTGTGCCGTCTCGCAAGCAGACCCGCTCACTGCCATTGACATCCT
CACCACCTGTGTCAGCAGACATCCAACGGCAGAGGTTCTTGCAGTGCACCGAGAAGGATCTGATTCCGTACCTGGA
GAAGCTAAGTGACAGCACGCTCAAGGAAACGCTGCTAAATGGGGTGGGCTACCTGCATGAGGGGCTCAGCCCCAT
GGAGCGACGCCTGGTGGAGCAGCTCTCAGCTCAGGGGCTATCCAGGTGGTGGTGGCTTCTCGGAGTCTCTGCTG
GGGCATGAACGTGGCTGCCACCTGGTAATCATCATGGATACCCAGTACTACAATGGCAAGATCCACGCCATATGT
GGATTACCCCATCTATGACGTGCTTCAGATGGTGGGCCACGCCAACCGCCTTTGCAAGGACGATGAGGGGCGCTG
TGTCAATCATGTGTGTCAGGGCTCCAAGAAGGATTTCTTCAAGAAGTTCTTATATGAGCCATTGCCAGTAGAATCTCA
CCTGGACCACTGTATGCATGACCACTTCAATGCTGAGATCGTACCAAGACCATTGAGAACAAGCAGGATGCTGT
GGACTACCTCACCTGGACCTTTCTGTACCGCCGCATGACACAGAACCCCAATTACTACAACCTGCAGGGCATCTC
CCATCGTCACTTGTTCGGACCACTTGTGAGAGCTGGTGGAGCAGACCCTGAGTGACCTGGAGCAGTCCAAGTGCAT
CAGCATCGAGGACGAGATGGACGTGGCGCCTCTGAACCTAGGCATGATCGCCGCTACTATTACATCAACTACAC
CACCATTGAGCTCTTCAGCATGTCCCTCAATGCCAAGACCAAGGTGCGAGGGCTTATCGAGATCATCTCCAATGC
AGCAGAGTATGAGAACATTCCCATCCGGCACCATGAAGACAATCTCCTGAGGCAGTTGGCTCAGAAGTCCCCCA
CAAGCTGAATAACCTTAAGTTCAATGATCCGCACGTCAAGACCAACCTGCTCCTGCAGGCTCACTTGTCTCGCAT
GCAGCTGAGTGCTGAGTTGCAGTCAGATACGGAGGAAATCCTTAGTAAGGCAATCCGGCTCATCCAGGCTCGGT
GGATGTCTTTCCAGCAATGGGTGGCTCAGCCCTGCTCTGGCAGCTATGGAAGTGGCCAGATGGTCACCCAAGC
CATGTGGTCCAAGGACTCATCCTGAAGCAGCTGCCACACTTCACCTCTGAGCATATCAACGTTGCACAGACAA
GGGAGTGGAGAGTGTTTTCGACATCATGGAGATGGAGGATGAAGAACGGAACCGGTGCTTCAGCTGACTGACAG
CCAGATTGCAGATGTGGCTCGCTTTTGTAAACCGCTACCTTAATATCGAACTATCTTATGAGGTGGTAGATAAGGA
CAGCATCCGAGTGGCGGGCCAGTTGTGGTGTGGTGCAGCTGGAGCGAGAGGAGGAAGTCACAGGCCCTGTCTAT
TGGGCTCTCTCTCCCGCAGAAACGTGAAGAGGGCTGGTGGGTGGTGATTGGAGATGCCAAGTCCAATAGCCTCAT
CTCCATCAAGAGCGTACCTTGCAGCAGAAGGCCAAGGTGAAGTTGGACTTTGTGGCCCCAGCCACTGGTGGCCA
CAACTACACTCTGTACTTCATGAGTGACGCTTACATGGGATGTGACCAGGAGTACAAATTCAGCGTGGATGTGAA
AGAAGCTGAGACAGACAGTGATTGAGATTGAGTCTGAGGCATTTACTTTTGGGTAAAGGAGAGTTGAGCCTGAA
TTAGGAATGTGTACATTGTAGGAATCCTGGTTGTGGGGACCAGGTCTGTGGGCCTCAGGTCTGGCCAGCCAGGGC
TGGTGTCTGTCCCGCCTACCTCCACTTCCTTTCCCTTGCTCACTCTGGATCCAGTGACAGCAGGTGTCTATGGGT
AAGCATAAATCATATATAGCATTTTCAGGCATGTTCTCTGGTAGTTCTTTTGTAGTCTGACATTCTAATAAAAAATAAT
TTGTAGAAACC

882/6881
FIGURE 821

MADVARSLOEYKANSNLVLQADRSIDRTRRDEPTGEVLSLVGKLEGTRMGDKAQRTPQMQEERRAKRRKRDR
EDRHDINKMKGYTLLSEGIDEMVGIIYKPKTKETRETYEVLLSFIQAALGDQPRDILCGAADEVLAFLKNEKLRD
KERRKEIDLLLGQTDTRYHVLVNLGKKITDYGGDKETIQNMDDNIDETYGVNVQFESDEEEGEDVDYGEVREEAS
DDMEGDEAVVRCITLSANLVASGELMSSKKKDLHPRDIDAFWLQRQLSRFYDDAIVSQKKADEVLEILKTASDDR
ECENQIVLLLGFNFTDFIKVLRQHRMILYCTLLASQAQSEAEKERIMGKMEADPELSKFLYQLHETEKEDLIREE
RSRRERVQRSMDDTLEMDLDQGGELAPRQVLDLEDLVFTQGSFHMANKRCQLPDGSRFRQRKGYEEVHVPAL
KPKPFGSEBQLLPVEKLPKYAQAGFEGFKTLNRIQSKLYRAALETDENLLCAPTGAGKTINVALMCMLEIGKHI
NMDGTINVDDFKIIYIAPMRSLVQEMVGSFGKRLATYGITVAELTGDHQLCKEEISATQIIIVCTPEKWDIIITRG
GERTYTQLVRLIILDEIHLHDDRGVLEALVARAIRNIEMTQEDVRLIGLSATLPNYEDVATFLRVDPAGKLFY
FDNSFRPVPLEQTYVGITEKKAIRKQIMNEIVYEKIMEHAGKNQVLVVFHSRKETGKTARAIRDMCLEKDTLGL
FLREGSASTEVLRTEAEQCKNLELKDLPYGFAIHAGMTRVDRTLVEDLFADKHIQVLVSTATLAWGVNLPALT
VVIKGTQVYSPEKGRWTELGAIDLQMLGRAGRPQYDTKGEGILITSHGELQYVLSLLNQQLPIESQMVSKLPDM
LNAEIVLGNVQNAKDAVNWLGYAYLYIRMLRSPTLYGISHDDLKGDPLLDQRRLDLVHTAALMLDKNNLVKYDKK
TGNFQVTELGRIASHYIITNDTVQTYNQLLKPTLSEIELFRVFSLSSEFKNITVREEEKELEQLKLLERVPIPVKE
SIEEPSAKINVLLQAFISQLKLEGFALMADMVYVTSAGRLMRAIFEIVLNRGWAQLTDKTLNLCKMIDKRMWQS
MCPLRQFRKLPEEVVKKIEKKNFPERLYDLNHNEIGELIRMPKMGKTIHKYVHLFPKLELSVHLQPIRSTLKV
ELTITPDFQWDEKVGSSAEAFWILVEDVDSEVILHHEYFLKAKYAQDEHLITFFVPVFEPLPPQYFIRVVS DRW
LSCETQLPVSFRHLILPEKYPPPTLELQLPVSALRNSAFESLYQDKFPFFNP IQTQVFNTVYNSDDNVFVGA
PTGSGKTICAEFAILRMLLQSSEGRVCYIIPMEALAEQVMDWYEFQDRNLKVVLLTGETSTDLLKLGKNI
ISTPEKWDILSRRWKQRKNVQINLFFVDEVHLIGGENGPVLEVICSRMRYISSQIERPIRIVALSSSLNAKDV
AHLGCSATSTFNHFNVRPVPLELHIQGFNISHTQIRLLSMAKPVYHAITKHSPKPVIVFVPSRKQTRLTAID
ILTTCAADIQRQRFHLCTEKDLIPYLEKLSDSLKETLLNGVGYLHEGLSPMERRIVEQLFSSGAIQVVVASRSL
CWGMNVAAHLVIMDTQYNGKIHAYVDYPIYDVLQMVGHANRPLQDDEGRVCIMCQGSKKDFFKKFLYEPLPVE
SHLDHCHMDHFNAEIVTKTIENKQDAVDYLTWTFLYRRMTQNPNYNLQGISHRHLSHLSELVEQTLSDLEQSK
CISIEDMDVAPLNLGMIAAYYIYNTTIELFSMSLNAKTKVRGLIEIISNAEYENIPIRHHEDNLLRQLAQKV
PHKLNNPKFNDPHVKTNLLQAHLSRMQLSAELQSDTEEILSKAIRLIQACVDVLSNGWLSPALAAMELAQMVT
QAMWSKDSYKQLPHFTSEHIKRCTDKGVESVFDIMEMEDEERNALLQLTDSQIADVARFCNRYPNIELSYEVVD
KDSIRSGGPVVVLVQLEREEVETGPVIAPLFPQKREEGWVVGDAKSNLSISIKRLTLQQAQKVKLDFVAPATG
AHNYTLYFMSDAYMGCDQEQYKFSVDVKEAETDSDSD

883/6881
FIGURE 822

GTTTCCATCCCTCCTCTGGAGGCTTTTGAAGTCACCGGGAGACAGATGTGCTCTGTGGCAGGCAGGGGAGTGGGG
GTGCTCAGGCACCTTGGGAGGTCTGGGAGCTCCTCAGTGTGACCGCTGGGACCCACCAGGACTTTTTCTTTGTCA
GAAGCCTTTGGTTGCTTTGCTGCTCTGCATGTGTCACTGTGGAGGGGCAATAGAGCAAGGCCTTACATGGCATGG
TCATTTCTCGGGCCAGGAGGCTTAGAGGCCTGCCCCTGGCGCTCAAGTATTGAACCAGAACCATGGGGTGGCAC
TGAGCCTCCTCACCACATCATGATAAATAACGGGGACATTACAGAGCAGGCACTGTTTCCTCAGTCCATGGCT
GAGTACATCACCGGTGTTTTCTCTCTTATTCCCTCCCATCAAGCCTAAAAGGAATCTCTATTGGAGATACTGCCAT
TAGTGTTCCTTTTATAGGTGAGGAACAGGCGATAGAGGGTTCCCCAGTTGAACCAACTGATAAATAGTAGAACT
TGGATTTTAATTTCAGTCTTGATGCCAGGGATAAGGCTCTTACTTTCTACCTTAGGCTATTTCTAGGAAACGCAGG
AGAGTGTTGAAGGGGCAGAGAAAGGGATCCAGTTCCTTTCTGTCCCGCATCCTAGTCCCTGAGAAGCAAAGAAGA
ATGTGTGGCTTCTTTTGCTTTGCTTTTGTGTGTCATCCACACATCTCCAGGGGACCTGGGCTCTTGATCTTGGCC
TCTTCCCCTTTAAGTGTAAAGTGGGAGCAGGTAAGGGGGTACAGTAGGGCTGGCCTGGAGTTAGAGGCTTGGATG
CCTTAGCTCCTCTGTCTGCACTCCAGAACTGCCTGACTTCATTTTCGTATGTTGTCTTTGTTTGGACAATTGATC
CATGTCCCAGTCCGCTCTTCTTCTTCTTCTTGATACTTACACTGCTTCTTCTGTTGGTTTTCCAGTGTTTAACACT
GTATACAACAGTGACGACAACGTGTTTGTGGGGGGCCCCACGGGCAGCGGGAAGACTATTTGTGCAGAGTTTGGC
ATCCTGCGAATGCTGCTGCAGAGCTCGGAGGGGCGCTGTGTGTACATCACCCCATGGAGGCCCTGGCAGAGCAG
GTATGACGTGGCGCTGTGTCATGTGAATTTCCCAAGAAGCATTTTCATCTGTGATTCCGTATGAAGGCTTTCTAAG
CCCTGAAATTTGCAGGGTCATTTCTCAGTTTGTGTATTAAAGAAAAGCTGCCCCAGCCAAGCGTGGTGGCTCAC
GCCTGTAATCCCAGCACTTTGGGAGGCCGAGGCGGGCAGATCTCCGGAGATCAGGAGTTTCGAGACCAGCCTGGCC
AACATGGTGAAACCCTGTCTCTACTAAAAATACAGAAATTAGCTGGGCGTGGTGGTGTGCGCCTGTAATCCCAGC
TACTTGGAAGGCTGAGGCAGGAGAATCGCTTGAACCCGGGAGGCGGAGGTTGCAGTGAGCCAAGTTCGCACCACT
GCACTCCAGCCTGGGCAACAAGAGCGAGACTTCATCTCA

884/6881
FIGURE 823

GC GGAAGGCGCGGGAGCTTGCGTGCTGCTGGGCTGAGCTGTCTGTCTCGTTTCTGTCCGCGCGCCCTGCATCCC
GGCCCCGGGCGCCCCGCTGGAGGTCGCGGAGGAGCCACAGGGCTGACTGGTCTGTGCCCCGGGCCCCAGGAGTGCCT
GGTGTAGCAGTCGCGGAGCCATCCC GGCGTCTGCTGCCATGACCGACTCTCCCCCTCAGAGGAGACTCTTCTCTCAG
CGGTGGCTGCAGAGACAGATGAGCGGCGGGCTCCTGGCCGCGGGACCGTGAGACGGGTTCGTGGCCGGCCATTTAG
GGGGACGCTGCGACCACCGCTGCGCCCCCTCCGACTGGTTCCCTTGGGCCCCGGAAGCTCGCGGCGGGGCCCTGCG
GGAGGCGGCATGCTCCCGCGGAGGCTGCTGGCCGCTGGCTGGCGGGGACGCGGGGCGGGGGCCCTGTGCGCGCTT
CTGGCCAATCAGTGCCGCTTCGTACAGGGCTGCGCGTGCGGCGCGCGCAGCAGATCGCGCAGCTCTACGGCCGC
CTCTACTCCGAGAGCTCACGCCGCTTCTCCTCGGCCGCTCTGGCGCCGGCTGCACGGCCGTCTTGCCATGCC
TCTGCCTTGATGGCGGCGTTAGCCGGCGTCTTCGTTTGGGACGAGGAGAGGATCCAGGAGGAGGAGTTGCAGAGA
TCTATTAATGAGATGAAGCGGTTGGAAGAAATGTCAAATATGTTTCAGAGCTCTGGAGTCCAGCACCACCTCCA
GAACCAAAAGCCCAAACAGAAGGGAATGAAGATTGAGAGGGCAAGAGCAACGTTGGGAAATGGTGATGGATAAG
AAACACTTTTAAGCTGTGGCGGCGCCCAATTACAGGCACCCACCTTTACCAGTACCGAGTTTTTGGAACTACACA
GATGTGACACCTCGGCAGTTCTTCAATGTTTACGTGGACACAGAGTATAGAAAAAATGGGATGCCCTGGTAATC
AAGCTGGAGGTGATTGAGAGGGATGTGGTTAGTGGTTCCGAGGTTCTTCACTGGGTAAACCATTTTCCCTATCCA
ATGTACTCACGGGATTATGTTTTATGTTCCGGCGGTATAGTGTGGATCAGGAAAACAACATGATGGTGTGGTGTGCG
CGTGCTGTGGAGCATCCGAGTGTGCCAGAGTCTCCAGAATTGCTCAGGGTCAGATCATATGAATCCCAAATGGTT
ATCCGTCCCCACAAGTCATTTGATGAGAATGGCTTTGACTACTTACTAACATACAGTGACAATCCCCAAACGGTG
TTTCTCTGCTACTGTGTTAGTTGGATGGTTTTCCAGTGGCATGCCAGATTTCTTGAGAAGCTGCACATGGCCACT
CTGAAAGCCAAGATATGGAGATTAAAGTAAAGGACTACATCTCAGCTAAGCCTCTGGAAATGAGTAGTGAAGCC
AAGGCCACCAGCCAGTCTCTGAGCGAAAGAACGAGGGCAGCTGTGGCCCTGCTCGGATTGAGTATGCTTGACAG
GCTTTTGGGATAAGAAGGGACAAGGTGCTTCTAGCCCTGTCTCAGTCCGTATCACTCTGCTGTAGAAGGGGGACA
TGCCACATGTATTAGAAGGCATCTGCTGTAACTTCCAGTGCAAGATAATTCAATAACTGATGTCCCATTTCATTC
AGAGCCCTTATTGCTCTTATCAAAACAGAAGAAGGTACATTTGTGGGAGTGTGTATATTCTCAGGCCAACTG
TTTTGAAATTCCGTATCTCACTGAGCTAATCTGGAACAAACCTCTCACCTCAGGCCAGAAGGGGATGACCTCCAT
TTGCTTCTCTGAGTAGTTTCTCTGCTGACATTCCAAATCCCACCATCGATTGTGCAGCGCTTTGGATTTCCTTC
AGTTCTCCAGGTCCACCTGGAAAGTATAGTTGGCCAGTTGAGTCTCTCAAATGAGGGGCTACTGGGAGTGCCTCT
GGTAACAATCATGATGTGAATGGGTGTGAACGATACTTGGCTATGTTAAGTGCCTTGTCCGACCTTGCTTTTAT
CTCTAGAGACATGAAGTTATTATTAATTTTTTTTTTTTAAAGTAGAGATGGAGTTTCACTCTGTTTCCCAGGT
GGTCTTGAACCTCCTGGGCCATGCCCTGGCCAGGGACATGAATTTGTACAAAGAAATTTCCCTCCCTGCACAA
TATCACCCATTGACTACCTTATCCAAAGCAAGTTTCTGTGAATCGGCCAGTTCTTCTATATTCACTGGATCAT
TGCCCTCCTTCTAACCTTCCCCATTTACCAAGAACTGGGAGACTAATCCTTTTAGATAGTAGCTTTTGTATGC
TCAAAACATCACATTTAAATTTAGTTTAAAAATTTTTTAACTTTTGTGTCAAATAGGAGTTGAGGAATTGAGCAG
GATTCTACCCTAGTCCGATTGTATAGAAAACACCATTTTGATTGAGGTATTATTTTTCATATTTCAGGTTTGACT
TGTTCTTTTCAGAAGGCTAAAGTCAGAGGAATGGGGCTGGGCCACTCCCTTGAGCTCTCAGATCTACAGACAA
GCTGTGTGAATGCATAGATGTAATCTTGTCTCAAATACTAATACAGTGGAGATTGGGTTTATGTTACCATTAAGT
TCCTCTAAAAAGTTTTTCTTCTCTCTTCAGAGCCAAAATAAAAGTGAACCTACACTGTTTCAAGATAAGGTACAAAT
CTGATGCTGTGAGTTTGACCGAGCTGGTTTTGCTTATGGTCATGCTGCAATTGTTAGAATAATAGGGATCAAGT
TTTAAATCCTCCTCCTTCCCTTTTTTCTGGAGTCTTGAGGGCCAGAGTTTTTGTTTTTGTTTTGTTTTGTTTT
CCTGCTTGCTACTGTTTTGTGGTGTGAAAAGTGGTTTAAACCTGAGACTAACTTAAACACTTCCCTGACCTTCT
TGTTCCTGTTTCAATTTTTGTGCCAAGGAAGTAGCTGCCCCAGTGTATGTCTTGCCTTCTCCGCGTCATTGTTGGA
AGAGGAGAGATGCATCGAGCAGTCCCAGCTGCTTTTCACTTACTTCTTCTTTCCAGGACCTGACAGAAGTCA
GGGAAGAGTCCCTGGGTTATGTCCAACTTAGCACCTGCAATTGTTGGGATGTGGATGGATGTGTGCATAAGAGA
GAGAGAGAATATGTGTGTGTGTGTGCGTCTGCGAGCGCACACATGCACAAGTGCGAAGGAGTTGCGGTTGCG
TCCATGTTCTGACTTAGGGCAATTTGATTCTGCACTTGGGGTCTGTCTGTACAGTTACTCATGTGATTGTAATGA
TTTCACTCCTAACTGTGACATTTTATCAAATGTGTGAATAAATACATAAAGATTGGTAC

885/6881
FIGURE 824A

CGCGCTTAGGGATCCGGCCGTGGCCGAGCGCGCGGCCGTAAGACCGCGGGTGACTAGCATGCAGATACCCATGCT
CTGACTTTCTGCCCCCTCCACTGACATGGCCACCGGGTGGGAGAGGGACTTCCAGACTTCAGCTCGACGCATG
GGCACCTCGTGCTCTTCCAGCTTTCAGTGCATGAACGGGAGCTGGACCTGGTTTTCTGGATCATAGCTATGCC
AAGCCTTGGAGTGCCACCCAGATGCCAGTAGTGCCCGCCCCACCCGCATGCTCTTGTCACTCCCCGGCGGCAG
CACGAAAGTACCATTGAATCAGACGTCCCAATAGATGTGGAGACGGTCACATCAACGCCTATGCCACTCTATGAC
AATCAGAAGGCACGCAGCGTGATGAATGAGTGTGAACGGCATGTCATCTTGGCAGGACTGATGCAGATGCCCT
CCTCCACCAGAGGACTGGGAGGAGCATGTCAACAGGACTGGCTGGACAATGGCCAGAACAAAGCTATTCAACAAG
ATCCTCAAAGCCCTGCAGTCTGACCGGCTTGCCCGCTTGGCCAACGAAGGGGCTTGTAAATGAGCCAGTGTGCGC
CGTGTGGCTGTGGACAAGTGTGCAAGGAGAGTGCGGCAGGCTCTGGCAAGTGTGAGCTGGGATACCAAGTGTATC
CAGTGGCTGCACACCACCCTTGTGGAGACCTTGAGTCTGCCCATGCTGGCAGCCTACCTGGATGCTTGTGAGACG
CTGAAGGGGAAGATCCCAACCTTGATTGACCGGATGCTTGTGTCTATCCAACACAAAGACTGGGGCTGCAGGAGCT
GAGGCCTTGTCTCTCTACTGAAGAGGCCCTGGGACCCTGCTGTGGGTGTGCTTTCTATAACAAACCAAGCAAA
CTCCCTGGCTCTCCGCTGATTCTCATCGCCTCCTCTGGTCCCTCCAGCTCTGTGTTTTCCACTTACGCCGCCAC
CGCTTCTGGCAATCTCAGCTGTCTGCTTGGGCAAGTTCATCCCTGTAGCCACCCATCTGCTGAACAATGGCAGT
GGGGTAGGAGTTCTACAGTGTCTCGAGCATATGATTGGGGCAGTGAGAAGCAAAGTGTGAGATTACAGCCAT
TCCCCACACAAACCCATTATCTTGATTGGCTGGAACACAGGAGCTTTGGTGGCCTGTCTGTGTGATGATGGAG
TATGTCACTGCAGTTGTCTGCCTTGGGTTTCTCTGCTTACTGTGGATGGCCCCAGAGGGGATGTAGATGATCCC
CTCTTGATATGAAGACTCCAGTCTCTTGTCTATTGGTCAGAAATCCCTTCAATGTACCCCTGAAGCCATGGAG
GACTTCCGGGAGAAGATTTCAGCTGAGAACAGCTTGGTGGTGGTTGGGGAGCTGATGACAATCTCAGAATAAGC
AAAGCAAAGAAGAAATCAGAAGGGTTGACTCAGAGCATGGTGGACAGATGATTACAGGATGAGATTGTGGACTTT
CTGACTGGAGTGTCACTCGTGTCTGAGGGTCACATGGGCTCTGAACTCGGGATCAGGATGCTGAGAAGAAGAG
AAGCCCCGCGATGTGGCCCGCAGAGACTTGGCCTTTGAAGTCCCCTGAGCGGGCAGTCGACCTGCCTCCCCAGCT
GCCAAGCTGCCCGCCTCACCTCAGGCTCAGAGGATCTCTCCAGTGTGTCCAGCAGCCCCACCTCCAGTCCCAAG
ACCAAAGTGACCACAGTGACCTCTGCCCAGAAGTCCAGTTCAGATTGGAAGTTCTCAGCTGCTGAAGAGACATGTG
CAGCGGACAGAAGCTGTGCTGACCCACAAACAAGCTCAAGCACAGTTTGCTGCTTTTCTGAAACAAAATATGCTG
GTGAGGAAAGCTCTTCTCCTGGCACCTCCTCTGCTCTTTTGTTCCTTTTCATCAGAACCACAGAGGAAGGA
GAGAAAGAGGATCTTAGGGTTTCAGTGAAGCGACACCATCCCTCGAGTCCCCTTCTGGCAGTAAGACCTCCAAA
CGACCGAAGATCAAGGTGTCCCTTATCTCCCAAGGGGACACAGCTGGAGGGCCTTGTGCTCCTTCCCAAGGAAGT
GCTCCAGAAGCTGCAGGTGGGAAGCCCATCCATGACACTGGGGCAGGCTTCAGCAGGGGCCAAGGAGCTCACA
GGACTTCTCACCACAGCCAAGTCCAGTTCTTCTGAAGGTGGAGTCTCAGCCAGCCAGTCCCTTCAGTGGTCTCC
AGCAGCACTGCACCCAGTGCCCTTGACACACTGCAGAGCCGCTGGTGGCCACATCTCCTGGCAGTCCCTCCCA
GGGGCCACATCAGCCAGCAGCCTCCTCCAAGGCTCAGCTTTCAGCTTGCAGGATATCAGCAGCAAGACCTCTGGC
CTTCCAGCAAATCCCTCCCCAGGACCAGCCCCACAGGCTTGAAGGTCCCCACCACCATTACTCTGACACTTCGTG
GCCAGCCGAGCAGGATCACTACACTGAGCCCTATGGGCTCAGGAGCAGCCCCATCCGAGGAGTCTCTTCCCAGG
TGCTGCCCTCCAGCTCACAGCGCCTGCCTCCAGCACCCTGAAGATGCTGTGTGATATGTCTCTTACCAAGTTG
GTGATGGCTGCCTCATGGTGGGCCCTGGACAGGTGTGTGGTCTGCTGAGCTGTCCACGTGTGGAAGACCTGTT
TAAGACAGTCATTTTTGCCTCTCCGCCAACTGTCTTCAGAGAAACCATTAGGTTAGGTGATACGGTGCCAGCAAG
GGAAGCACCATCGTCCAGGATCTGCAAATCTGGTTCTGGGAACCCAGACTCCTCAGCAGATCTGGCTGTACAT
GGATCAGAACCACCTTCTTCCCCCGCTTAAGCTGTGGTTGACCCAAGGGTCAGCATATAGGACTGCCTGTGCTCAT
TTAATGAAGGTGTTTTCTTTTGAAGTCTGTGCTACCCTCTGCGCCTAGTTGGGAGGAGACATCCATCTGGTCTG
GGATTTCCGGAGTTAGAATGGAAGCTCTTTGTCTAAAGACTGGAGTCATCTGGCCTGCCAACTGGTGGTTCAGA
GCCGGACGGGCTTGTTTTGGACATCACTGTTGCCTTCACTCAGCAGCCACGGGAGAGTGTCCCCATGCAACTCC
ACCTTAGAAACACGTCAGATACTGAGTAGCTTGTGACTCCTGGAACTTCTGGTTTTTGTAGTATCATAATG
AAGGCAAAGAGTAAGTGTCTCTTTTTCAGCCTCTTTGACTTACTCTAGATGTTGGGAGCAGTGGTTGCCAGG
TGAAACCTGGGCCCTTTGTCTTTTACCATGCTTTGGGCAGTTTCTGTATCCAGAGAGTCCGCAGGTTTCAGATA
AGCTGAAGAAGAGTAATAGAACAGCAAAGGAAGTGGCTGAAGGATGTGCTAGTAAGCCCTGTGGTTTTGTGCTTA
GGTCTCTGCTCTGCTACCCAAGGAAGTGGTGGTTTCAGCTGGAGATAAAAAGAAGAATTTGCCAAGTCAGAGAAGA
AACCCCAACCCCGAAAATCCTCTGTCTCCAGTCTCTGGAGGTGAAGCAGGGACAATAAGCTAAGGTAGTATCTT

886/6881

FIGURE 824B

GGCCATCCCAGGAACTTGTGGCATTAGGACGATGAAGGCCATGCTTCAGTGTTTTTCGTTTCTATTTTCATGAGAC
TTTTTGTCTTCCTGCTTACAAGTGGGAAGATGATTGACAGTGACTCTACTATGCAGGGCTGTTGGTACCAACCTG
AGCCCTATAGGTGGCAGTCCCTGGAGAAGTGGTCACAGAAGATGGAGCTCTGATCCCTGCTTACCTCTTACAA
CACTTGTGTGCAAAGATAGTTTTAGATTTGGTTTAGAAGCTATCCTCCAGAACAGGCTCCCATACTTAGAATGTT
TCTAGTTAAGGTAATAAATTAGGCAACCCAAGTGTGACTCCACTCAAGTGTCTTTTTCTGTAGGCAGGAAGGGCC
CACAACATGGCTTAAATGTAGTCCATGGTTCTGGCCACAGTACAGTGTGTATCTATACCAGGTCACCTGTGTT
CAATCTGGGAGCCTTCCTGGCCAGTCTGAGTGGCAGCCAGAAGGGAGCTCATAGTGTCTAGGAGTCTCAGGCAAG
GTAGGTCAGGGTACTGTGGGCAGGGGGGATGTGTGTGATAGGAGAGGGTACCCTAAACCCCATACCTTCCCTCCC
TGACCTGAAAAGCTGATCTCAACAGGGATTACACAGAATTAGGCTGTGTTTTTGCATTAGCTGGTAGGTGACTT
TCTCAAATTTCTTAAATTCAGAAAGTATTTAGTAAACTTGAGGAAGGTATGAAATCTGGAGGAGGCATCCAGGAC
CCAGGGGTTTTGATAGCTTTACAGGTAGGATCATACCACACCAAAAGAGCAGTGGACAATAAGACTATATGAGCTA
TATGAAGCTTTTAGGAATCATTTAGGACAGACAGAGCCCTAAACAACCCATTATGACTTAAGTTGTTGGCTCAG
TGATGCTGGGGACAAAGAAAACTAACAAGCCGACCTGCCTTTATGATAAATTCTAGTGTGCTTACAAGGGATG
ACTTCTGAGGTGTGATCTGTCCACCTTGAAGAACTCCACAACCTGAAGAAGGGGAGCTGTGAGAACGTGGATTGT
TCTACAACCTTGCACAGGGTAACAGAGGAAGTGGCTGAGGCCTAGAGTCACGTTTTCCAGTTCCTTCGCAAACTA
TATTTCTTGGAACGCGAAAGGAAGCTTTACCTATTTTATAGAAGACCTGGAATCCATAACCTCAGAAGGCAATAT
TATTGATAGAAAATGTGGAAGGATCAGGAAGTTCTTAGATTCTTGGATGACAGATGCATGTTGATGCCCTATGGA
GATGTCCTTGTGTTTGGAGGTCACTGAGGTAGGAAGACCTGTCTACTCTTGGTTTCACCACTAGAACAGTCTTGG
GCTGGATGGGTTATAGAGCTGAGCGGCTGTGATGGTTCTGTTTTTACATTAAACAAAACAATTAACCAACCAAA
AAC

FIGURE 825

[illegible]

888/6881
FIGURE 826

MYLSAICEYQLEEIQRVFEGPYKEYHEEAQKWDRYTDVPSPRPGSCINNWHRHGYTSSLELPDNILNFVKKHP
LMEEQVGPRWSRPLLVKKGTFNTHLVADRVTLGDLGATYTVLFIDTGDGWLLKAVSLGPVHLEELQLFDQEPMR
SLVLSQSKKLLFAGSRSQVLVQLPVADCMKYRSCADCVLARDPYCAWSVNTSRCVAVGGHSGSLLIQHVMTSDTSG
ICNLRGSKKVRPTPKNITVVAGTDLVLPCHLSSNLAHARWTFGGRDLPAEQPGSFLYDARLQALVVMQAQPRHAG
AYHCFSEEQGARLAAEGYLVAVVAGPSVTLEARAPLENLGLVWLAVVALGAVCLVLLLLVLSLRRRLREELEKGA
KATERTLVYPLELPKEPTSPFFRPCPEPDEKLWDPVGYYYSDGSLKIVPGHARCQPGGGPPSPPPGIPGQPLPSP
TRLHLGGGRNSNANGYVRLQLGGEDRGGLGHPLPELADELRRKLQQRQPLPDSNPEESSV

889/6881

FIGURE 827

CGGAAGTTTTGCTGCTAGTCGCGGACGCAATGGCTTCAAGGTTACTTCGCGGAGCTGGAACGCTGGCCGCGCAGG
CCCTGAGGGCTCGCGGCCCCAGTGGCGCGGCCGCGATGCGCTCCATGGCATCTGGAGGTGGTGTTCCTACTGATG
AAGAGCAGGCGACTGGGTTGGAGAGGGAGATCATGCTGGCTGCAAAGAAGGGACTGGACCCATACAATGTACTGG
CCCCAAAGGGAGCTTCAGGCACCAGGGAAGACCCTAATTTAGTCCCCTCCATCTCCAACAAGAGAATAGTAGGCT
GCATCTGTGAAGAGGACAATACCAGCGTCGTCTGGTTTTGGCTGCACAAAGGCGAGGCCAGCGATGCCCCCGCT
GTGGAGCCCATTACAAGCTGGTGCCCCAGCAGCTGGCACACTAGGCACCTGCACTAAATTACTCAAAATGTGCTG
TAAAGTTCTTCTTTCCAGTAAAGACTAGCCATTGCATTGGCTCCTTCTCCATAAAAAAAAAAAAAAAAAA

890/6881
FIGURE 828

MASRLRGAGTLAAQALRARGP SGAAAMRSMASGGGVPTDEEQATGLEREIMLAAKKGLDPYNVLAPKGASGTRE
DPNLVPSISNKRIVGCICEEDNTSVVFWLHKGEAQRCPRCGAHYKLVPPQLAH

891/6881
FIGURE 829

TTGCAGCCGCCGGCAGCTACTGCAAGGCAAAAGCCGGAGTGGACGTGTCTTTTGAAACTGCTGCTCTTTCACTTC
TCAGGCGTCACCGAGAGCTCAGCACCCAGGCTGAACTCTGTACCATTTGGAAGAATGGAAGCTGATGCATCTGTT
GACATGTTTTCCAAAGTCCTGGAGCATCAGCTGCTTCAGACTACCAAAGTGGTGAAGAACATTTGGATTCTGAA
ATTCAAAAAGTGGATCAGATGGATGAGGATGAATTGGAACGCCTTAAAGAAAAGAGACTCCAGGCACTAAGGAAA
GCTCAACAGCAGAAACAAGAATGGCTTTCTAAAGGACATGGGAATACAGAGAAATCCCTAGTGAAGAGACTTT
TTTCAAGAAGTCAAGGAGAGTGAAAATGTGGTTTGCCATTTCTACAGAGACTCCACATTAGGTGTAAAATACTA
GACAGACATCTGGCAATATTGTCCAAGAAACACCTCGAGACCAAATTTTTGAAGCTGAATGTGGAAAAAGCACCT
TTCCTTTGTGAGAGACTGCATATCAAAGTCATTCCCACACTAGCACTGCTAAAAGATGGGAAAACACAAGATTAT
GTTGTTGGGTTTACTGACCTAGGAAATACAGATGACTTCACCACAGAACTTTAGAATGGAGGCTCGGTTCTTCT
GACATTCTTAATTACAGTGGAAATTTAATGGAGCCACCATTTCAGAACCAAAAGAAATTTGGAACAACTTCACA
AAGCTGGAAAAGAAACTATCCGAGGAAAGAAATATGATTGAGACTCTGATGATGATGATGAGCTCAATAATTCTT
TGTAATTTGCTTTTTTTTTTCTGCTTCAGATTTAAATGTGTTTTTAAATTTCTATTAATGTCTATACATTGGTCA
CCTAAATACTCATATTCTCGAGTTTATACAGTTGTATCACATCGAAAAGTGTCTTTACTGTTTTCTGTGTGGCC
ATCATGTTTAAAGTTGAGGAAAAGTCAAGTTCTTAAATTATCTGGGAAGGGTCTGGATTCTCTATTTTTGAGATTGA
CTTTATCACAATATGATTCTTACATCTTTATACCATTTACAATTGTGTTTTAGATCTACAGAGTTAGAAATTCGA
AACTATTCCAGGACTAATTCTTAATCGGCATTATTTATACAAGAGGTCAAGTAACATTTACTAGCGCAATACTG
CACTTGTAATGAATTATAAACGCTCTTCTGGAATATATTTAAATAACCATTAAAGAACTGCTTATTCATTCTGG
ACACTGCATGTTGATGTTGAATCACTGATGCCAGCAGAAAGCTATTTTGATTGTGAACATACTGCCTTATTTA
AAGGGTCCTGATTGCTTGTATTTAAGACATTCATTAAGAAAGAAACCAGGAAACACTTTTGAAATAACAGCATAA
GGAACCTTCACTGCTCTGCTCAATAAAATACCTGTAAGTGAAGAAAAAAGAAAAAAGAAAAAAGAAAAAAGAAAAA

892/6881
FIGURE 830

MEADASVDMFSKVLEHQLLQTTKLVEEHL DSEIQKLDQMD EDELERLKEKRLQALRKAQQQKQEWLSKGHGEYRE
IPSERDFFQEVKESENVVCHF YRDSTFRCKILDRHLAILSKKHLET KFLKLNVEKAPFLCERLHIKVIPTLALLK
DGKTQDYVVGFTDLGNTDDFTTETLEWRLGSSDILNYSGNLMEPPFQNKKF GTNFTKLEKKTIRGKKYDSDD
D

893/6881
FIGURE 831A

CGCGGGTCTGTGGAGAGCCGGGTGCGAGCGGCGGCAGCACGAGGGGAAAAGAGCTGAGCGGAGACCAAAGTCAGC
CGGGAGACAGTGGGTCTGTGAGAGACCGAATAGAGGGGCTGGGGCCACGAGCGCCATTGACAAGCAATGGGGAAG
AAACAGAAAAACAAGAGCGAAGACAGCACCAAGGATGACATTGATCTTGATGCCTTGGCTGCAGAAATAGAAGGA
GCTGGTGTCTGCCAAAGAACAGGAGCCTCAAAAGTCAAAGGGAAAAAGAAAAAAGAGAAAAAAGCAGGACTTT
GATGAAGATGATATCCTGAAAGAACTGGAAGAATTGTCTTTGGAAGCTCAAGGCATCAAAGCTGACAGAGAACT
GTTGCAGTGAAGCCAACAGAAAACAATGAAGAGGAATTCACCTCAAAGATAAAAAAAGAAAGGACAGAAGGGC
AAAAACAGAGTTTGTATGATAATGATAGCGAAGAATTGGAAGATAAAGATTCAAATCAAAAAAGACTGCAAAA
CCGAAAGTGGAATGTACTCTGGGAGTGATGATGATGATGATGATTTTAACTTCTTAAAAAGCTAAAGGAAAA
GCTCAAAAATCAAATAAGAAGTGGGATGGGTGAGAGGAGGATGAGGATAACAGTAAAAAATTAAGAGCGTTCA
AGAATAAATTCTTCTGGTGAAAGTGGTGATGAATCAGATGAATTTTGAATCTAGAAAAGGACAGAAAAAAT
CAGAAAAACAAGCCAGGTCTAACATAGAAAGTGGGAATGAAGATGATGACGCCTCCTTCAAATTAAGACAGTG
GCCCCAAAAGAAGGCAGAAAAGAAGGAGCGCGAGAGAAAAAAGCGAGATGAAGAAAAAGCGAAACTGCGGAAGCTG
AAAGAAAAAGAAGAGTTAGAACAGGTAAAAAGGATCAGAGTAAACAAAAGGAATCTCAAAGGAAATTTGAAGAA
GAACTGTAAATCCAAAGTGACTGTTGATACTGGAGTAATTCCTGCCTCTGAAGAGAAAGCAGAGACTCCCA
GCTGCAGAAGATGACAATGAAGGAGACAAAAAGAAGAAAGATAAGAGAAAAAGAAAGGAGAAAAAGGAAGAAAA
GAGAAAGAGAAGAAAAAGGACCTAGCAAAGCCACTGTTAAAGCTATGCAAGAAGCTCTGGCTAAGCTTAAAGAG
GAAGAAGAAAGACAGAAGAGAGAAGAGGAAGAACGTATAAACCGGCTTGAAGAATTAGAAGCCAAGCGTAAAGAA
GAGGAACGATTGGAACAAGAAAAAGAGAAAGGAAAAAGCAAAAAGAAAAAGAAAGAAAGAACGCTTGAAAAA
GAAGGGAAACTTTTAACTAAATCCAGAGAGAAGCCAGAGCCAGAGCCGAAGCTACTCTTAACTGCTACAAGCT
CAGGGTGTGTAAGTGCCATCAAAGACTCTTTGCCAAAGAAGAGGCCAATTTATGAAGATAAAAAAGAGGAAAAA
ATACCACAGCAGCTAGAAAGTAAAGAAGTGCTGTAATCAATGGAATTATGTGCTGCTGTAGAAGTTATGGAACAA
GGAGTACCAGAAAAAGGAGAGACACCACCTCCTGTTGAACCAGAAGAAGAAGATACTGAGGATGCTGGATTG
GATGATTGGGAAGCTATGGCCAGTGATGAGGAGACAGAAAAAGTAGAAGGAAACAAAGTTTCATATAGAAGTAAAA
GAAACCCCTGAAGAGGAGGAGGAGGAGGAAGAAGAGGAAGAAGAAGATGAAGAAAGTGAAGAAGAGGAGGAAGAG
GAGGGAGAAAGTGAAGGCAGTGAAGGTGATGAGGAAGATGAAAAGGTGTCAGATGAGAAGGATTGAGGGAAGACA
TTAGATAAAAAGCCAAAGTAAAGAAATGAGCTCAGATTCTGAATATGACTCTGATGATGATCGGACTAAAGAAGAA
AGGGCTTATGACAAAGCAAAACGGAGGATTGAGAAACGGCGACTTGAACATAGTAAAAATGTAAACACCGAAAAAG
CTAAGAGCCCCCTATTATCTGCGTACTTGGGCATGTGGACACAGGGAAGACAAAAATTCTAGATAAGCTCCGTCAC
ACACATGTACAAGATGGTGAAGCAGGTGGTATCACACAACAAATTGGGGCCACCAATGTTCTCTTGAAGCTATT
AATGAACAGACTAAGATGATTAAAAATTTTGATAGAGAGAATGTACGGATTCCAGGAATGCTAATTATTGATACT
CCTGGGCATGAATCTTTCAGTAATCTGAGAAATAGAGGAAGCTCTCTTTGTGACATTGCCATTTTAGTTGTTGAT
ATTATGCATGGTTTGGAGCCCCAGACAATTGAGTCTATCAACCTTCTCAAATCTAAAAATGTCCCTTCATTGTT
GCACTCAATAAGATTGATAGGTTATATGATTGAAAAAGAGTCTGACTCTGATGTGGCTGCTACTTTAAAGAAG
CAGAAAAAGAATACAAAGATGAATTTGAGGAGCGAGCAAAGGCTATTATTGTAGAATTTGCACAGCAGGGTTTG
AATGCTGCTTTGTTTTATGAGAATAAAGATCCCCGACTTTTGTGCTTTTGGTACCTACCTCTGCACATACTGGT
GATGGCATGGGAAGTCTGATCTACCTTCTTGTAGAGTTAACTCAGACCATGTTGAGCAAGAGACTTGCACACTGT
GAAGAGCTGAGAGCACAGGTGATGGAGGTTAAAGCTCTCCCGGGATGGGCACCACTATAGATGTCATCTTGATC
AATGGGCGTTTGAAGGAAGGAGATACAATCATTGTTCTGGAGTAGAAGGGCCATTGTAACCTCAGATTGAGGC
CTCCTGTTACCTCCTCCTATGAAGGAATTACGAGTGAAGAACCAGTATGAAAAGCATAAAGAAGTAGAAGCAGCT
CAGGGGGTAAAGATTCTTGGAAAAGACCTGGAGAAAACATTGGCTGGTTTACCCCTCCTTGTGGCTTATAAAGAA
GATGAAATCCCTGTTCTTAAAGATGAATTGATCCATGAGTTAAAGCAGACACTAAATGCTATCAAATTAGAAGAA
AAAGGAGTCTATGTCCAGGCATCTACACTGGGTTCTTTGGAAGCTCTACTGGAATTTCTGAAAACATCAGAAGTG
CCCTATGCAGGAATTAACATTGGCCAGTGATGATAAAAAAGATGTTATGAAGGCTTCAGTGATGTTGGAACATGAC
CCTCAGTATGCAGTAATTTTGGCCTTCGATGTGAGAATTGAACGAGATGCACAAGAAATGGCTGATAGTTTAGGA
GTTAGAATTTTGTAGTGAGAAATTTTATCATTATTTTATGATGCCTTTACAAAATATAGACAAGACTACAAGAAA
CAGAAACAAGAAGAATTTAAGCACATAGCAGTATTTCCCTGCAAGATAAAAAATCCTCCCTCAGTACATTTTAAAT
TCTCGAGATCCGATAGTGATGGGGGTGACGGTGGAAGCAGGTGAGTGAACAGGGGACACCCATGTGTGTCCCA
AGCAAAAATTTTGTGACATCGGAATAGTAACAAGTATTGAAATAAACCATAAACAAGTGGATGTTGCAAAAAA

894/6881

FIGURE 831B

GGACAAGAAGTTTGTGTAAAAATAGAACCTATCCCTGGTGAGTCACCCAAAATGTTTGGAAGACATTTTGAAGCT
ACAGATATTCTTGTTAGTAAGATCAGCCGGCAGTCCATTGATGCACTCAAAGACTGGTTCAGAGATGAAATGCAG
AAGAGTGACTGGCAGCTTATTGTGGAGCTGAAGAAAGTATTTGAAATCATCTAATTTTTTACATGGAGCAGGAA
CTGGAGTAAATGCAATACTGTGTTGTAATATCCCAACAAAATCAGACAAAAATGGAACAGACGTATTTGGACA
CTGATGGACTTAAGTATGGAAGGAAGAAAAATAGGTGTATAAAATGTTTTCCATGAGAAACCAAGAACTTACAC
TGGTTTGACAGTGGTCAGTTACATGTCCCCACAGTTCCAATGTGCCTGTTCACTCACCTCTCCCTTCCCCAACCC
TTCTCTACTTGGCTGCTGTTTTAAAGTTTGCCCTTCCCCAAATTTGGATTTTTATTACAGATCTAAAGCTCTTTC
GATTTTATACTGATTAAATCAGTACTGCAGTATTTGATTAAAC

895/6881

FIGURE 832

CTTCCTTTCCAACCTTGGACGCTGCAGAAATGGCTCCCGCAAAGAAGGGTGGCGAGAAGAAAAAGGGCCGTTCTGCC
ATCAACGAAGTGGTAACCCGAGAATACACCATCAACATTACAAGCGCATCCATGGAGTGGGCTTCAAGAAGCGT
GCACCTCGGGCACTCAAAGAGATTGGGAAATTTGCCATGAAGGAGATGGGAACCTCAGATGTGCGCATTGACACC
AGGCTCAACAAAGCTGTCTGGGCCAAAGGAATAAGGAATGTGCCATACCGAATCCGTGTGCGGCTGTCCAGAAAA
CGTAATGAGGATGAAGATTCACCAAATAAGCTATATACTTTGGTTACCTATGTACCTGTTACCACTTCAAAAAT
CTACAGACAGTCAATGTGGATGAGAACTAATCGCTGATCGTCAGATCAAATAAAGTTATAAAATTGC

896/6881
FIGURE 833

MAPAKKGGEKKKGRSAINVVTREYTTINIHKRIHGVGFKKRAPRALKEIRKFAMKEMGTPDVRIIDTRLNKAVWAK
GIRNVPIRIRVRLSRKRNEDEDSPNKLYTLVTYVPVITTFKNLQTVNVNEN

897/6881
FIGURE 834

ATGCCCCGGCGGAGGGGCGAGCGCGGCGTCTGGCCGGCTTCTACCGCCGCGGAGCAAAGAGGGTCCCGGGAAGCG
GCAGGGTCGGCGTCCAGGAGCGGGCTTCGGGGGCTCCGGCGGCGGCAGAGGCGGAGCAAGCGGGCCCCGGGTCCGGG
AGCGGAGGCCCTGGGGGCCCCGCGGGCAGGATGAGCTTGACCCCGAAGGAGCTCTCGAGCCTGCTGAGCATCATA
TCGGAGGAGGCGGGCGGCGGCAGCACCTTCGAGGGCCCTGTCCACCGCCTTCCACCACTACTTCAGCAAGGCCGAC
CACTTCCGCTGGGCTCGGTGCTCGTCATGCTGCTCCAGCAGCCCGACCTGCTGCCTAGCGCGGCGCAGCGCCTC
ACGGCGCTCTACCTGCTCTGGGAGATGTACCGCACCGAGCCGCTGGCCGCCAACCCTTCGCCGCCAGCTTCGCG
CACCTGCTCAACCCCGCGCCGCCCGCCGCGGCGGCCAGGAACCCGACCGCCCTCCGCTCTCAGGATTTTACCT
CCTATAACTCCACCAGAAAAGTTTTTCTTTCCAGCTGATGTGGCACCCCCACGGGAACCTCTTCAAAAAGACG
CCTCGCCAGATTGCACTGATGGACGTTGGAAACATGGGCCAGTCTGTGGACATTAGTGGGCTTCAGTTAGCCTTG
GCCGAACGCCAATCTGAATTGCCAACGCAAAGCAAAGCGAGCTTCCCCAGTATTCTCAGTGACCCAGACCCGGAT
TCTTCTAATTCTGGATTTGACAGCTCAGTTGCCTCTCAGATCACAGAAGCTTTAGTCAGCGGACCAAAGCCACCT
ATTGAAAGCCATTTTCGACCAGAGTTTATTGCTCCACCGCCTCCACTCCACATTTGTGAGGATGAACCTGCTTGG
CTAAACCCACGCGAGCCTGACCACGCGATCAGTGGGATAAATCGATGTGTGTTAAGAATAGCACTGGTGTGGAG
ATCAAACGAATAATGGCCAAAGCCTTCAAAGCCCTTATCCTCTCCCCAACAAACACAGCTACTTGGTGAGTTG
GAAAAAGACCCCAAACCTTGTCTACCATATTGGCCTCACCCAGCCAAACTTCTGACCTTGTGGAAAAACAACCT
TTAGTCGCTATAGAAATGTTGCTGAAATTAATGCAGTCAAGCCAGATCACTGAGTATTCTCTGTCCTGGTCAAT
ATGGACATGTCTTTACATTCAATGGAAGTTGTAAATCGACTAACTACAGCTGTTGATCTACCTCCTGAATTTATT
CACCTTTATATATCAAATTGCATCTCTACTTGTGAACAGATTAAGGATAAATATATGCAGAATCGGTTGGTGCGT
CTTGTGTGTGTGTTTCTCCAATCCTTGATCCGTAACAAAATTATTAATGTACAGGATTGTTTATAGAAGTGCAG
GCATTCTGTATTGAATTCAGTAGGATACGAGAAGCTGCTGGTCTTTTCCGGTTGTTGAAGACATTGGATACTGGG
GAAACACCTTCTGAGACCAAATGTCAAAATAA

898/6881

FIGURE 835

MPGGGASAASGRLLTAAEQRGSR EAAGSASRS GFGSGGGRGGASGPGSGSGGPGGPAGRMSLTPKELSSLLSII
SEEAGGGSTFEG LSTAFHHYFSKADHFRLG SVLVMLLQQPDLLPSAAQR L TALYLLWEMYRTEPLAANPFAASFA
HLLNPAPPARGGQEPDRPPLSGFLPPITPPEKFFLSQLMLAPPREL FKKTPRQIALMDVGNMGQSVDISGLQLAL
AERQSELPTQSKASFPSILSDPD PDSSNSGFDSSVASQITEALVSGPKPPIESHFRPEFIRPPPPLHICEDELA W
LNPTEPDHAIQWDKSMCVKNSTGVEIKRIMAKAFKSPLSSPQQTQLLGELEKDPKLVYHIGLTPAKLPDLVENNP
LVAIEMLLKLMQSSQITEYF SVLVNMDMSLHSMEVVNRLTTAVDLPPEFIHLYISNCISTCEQIKDKYMQNRLVR
LVCVFLQSLIRNKIINVQDLFIEVQAF CIEFSRIREAAGLFRLLKTLDTGETPSETKMSK

899/6881
FIGURE 836

AGGGTACGGGCCGGGACCGCCGACGCCGGGGCGGGGGCACGGCAACCGCGAGGCCTGGGGGCGCCCGCCCCCG
CGCCCCACGCCCGGTGCCAGCGAGCCGAGGCGTGATCTCTTATATGGTCAAATGACACGGCGGGGTTTCTCGA
GGGCGGGAGCTGCGCAGCGCTCCACTCGGCCGGCAGCGGAGCCGACGCCACCAGCCGCCCGCGCCCTCCAGCCCC
GTCCGGGAGTCCCCGGCCCGCTGCGGTGCCGTGAGTACCTCCAACCCCTGCGCCCCGGAGGGAGGCCGAGGGGC
TTAGCCACCAGGGCTCGGAAGTGGGGGCCGAATCCGGTGCGAGACCCAAGGAGAGGGGAGCAGAGCCGGAGTTGG
GGAGACTGTGGCTGAAAACGTGTCTTCTGGAGACTAGGCTGGCATTGACTTTGGGACGGAGTCTCGCTTG
TCGCCCAGGCTGGAGTGCAGTGGCAGATCTCAGCTCACTGCAAGCTCTACCTCTTGGTTCAGGCCATTCTCCTG
CCCCAGCCTCCAAGTAGCTGGGACTACAGGTTGCTGAAAAGCCAGGAGTCAAAATGACTGAGCGCTTTGACTGC
CACCATTGCAACGAATCTCTCTTTGGCAAGAAGTACATCCTGCGGGAGGAGAGCCCCCTACTGCGTGGTGTGCTTT
GAGACCCTGTTGCGCAACACCTGCGAGGAGTGTGGGAAGCCATCGGCTGTGACTGCAAGGACTTGTCTTACAAG
GACCGGCACCTGGCATGAAGCCTGTTTCCACTGCTCGCAGTGCAAACTCACTGGTGGACAAGCCCTTTGCTGCC
AAGGAGGACCAGCTGCTCTGTACAGACTGCTATTCCAACGAGTACTCATCCAAGTGCCAGGAATGCAAGAAGACC
ATCATGCCAGGTACCCGCAAGATGGAGTACAAGGGCAGCAGCTGGCATGAGACCTGCTTCATCTGCCACCGCTGC
CAGCAGCCAATTGGAACCAAGAGTTTCATCCCCAAGACAATCAGAATTTCTGTGTGCCCTGCTATGAGAAACAA
CATGCCATGCAGTGCGTTTCACTGCAAAAAGCCCATCACACGGGAGGGGTCACTTACCGGGAGCAGCCCTGGCAC
AAGGAGTGCTTCGTGTGCACCGCCTGCAGGAAGCAGCTGTCTGGGCAGCGCTTCACAGCTCGCGATGACTTTGCC
TACTGCCTGAAGTCTTCTGTGACTTGTATGCCAAGAAGTGTGCTGGGTGCACCAACCCCATCAGCGGACTTGGT
GGCACAATAATACATCTCCTTTGAGGAACGGCAGTGGCATAACGACTGCTTTAACTGTAAGAAGTGTCTCCCTCTCA
CTGGTGGGGCGTGGCTTCTCACAGAGAGGGACGACATCCTGTGCCCCGACTGTGGGAAAGACATCTGAATTCAA
CACAGAGAAGTTGCTGCTTGTGATCTCACACACAGATTTTATGTTTTCTTCTCACCCAGGCAATCTTGCCTTC
TGGTTTCTTCCAGCCACATTGAGACTTTCTTCTAGTGCTTTTCACTGATACTCACGTTTGCTTAAACCTTTAGT
GCTTTGTGATAGTTCACTCCAGGGAAGAGAAAACCTCGCCCTAGGCCCTAGGTGGGAAGATGGTTTGAATTTT
TGTAATCGAGTAAGGCACCCAAATGTAAAAATCCTTTTGAATGATGCCTTTATAAATCTTCTCTCACGTGCT
ATTTAAGTGCAATTAACATATGTCACGAACTTGAAAGTTTCTAAACTCAATAAGGTAATGACCAGTTGTTATTT
ACAGCTCTGTAACCTCCCGTTGCGTCAAGTCTAAACCAAGATTATGTGACTTGCAATAAAGTTATTCAGAACAGA
AAAAAAAAAAAAAAAAA

900/6881
FIGURE 837

MTERFDCHHCNESLFGKKYILREESPYCVVCFETLFANTCEECKPIGCDCKDLSYKDRHWHEACFHCSQCRNSL
VDKPFAAKEDQLLCDYCSNEYSSKCQECKKTIMPGTRKMEYKGSSWHETCFICHRCQQPIGTSFIPKDNQNF
VPCYEKQHAMQCVQCKKPITGGVITYREQPWKECFVCTACRKQLSGQRFRTARDDFAYCLNCFCDLYAKKCAGCT
NPISGLGGTKYISFEERQWHNDCFNCKKCSLSLVGRGFLTERDDILCPDCGKDI

901/6881
FIGURE 838

AGTCTTCTGGCCCCGAGAGGGAGAGGTGGGGATAGGACCTTCCTTTAGTCCCCTCCGTAGTGGAACGAAGGATTCTTTATGGTCATCTGIGGTGTTCAATTCGTAAAGAAGCATCTGCAGGGTAAGAAGGAAAAACGTCCACCTTGCAAA
TATATCCCAGGCACATGCCTCCTGAGAAGTGACCCCTCCTCCCTCCGCCTCCCCGGGCACGTCTGGGGCTTCTCCAGTCTCCCGCTCCTGGGACCAGGCAGAGATCCCGGCGTGGGCAGACCCCTGCCACCGCGCCCCAGGCCTCGTCC
GAAACTCGCGCCCGAGAGCCCACGCCGGAGGCGACGCTCCCGATAACTACTGTGCTCCCAAGACCCGGAGGGCACGAAGAAAGGAGCCCTGGCAAACAAAGGGTTGCTGAAAAGCCAGGAGTCAAAATGACTGAGCGCTTTGACTGCCAC
CATTGCAACGAATCTCTCTTTGGCAAGAAGTACATCCTGCGGGAGGAGAGCCCTACTGCGTGGTGTGCTTTGAGACCCGTTCGCCAACACCTGCGAGGAGTGTGGGAAGCCCATCGGCTGTGACTGCAAGGTACCTGCTGCGGGCCTC
TGCGGGTCCCCGGCGCTCCCTAGGAAACACGGGTTTCTCTGTGGTTTGTGCTGAAGCCCTGATCCACCACTGAA
GTCTGTCTCTTGAGCCCCCTCTGAGGGGCGTGGACCCTGCGTGTAGCCTTTCGGAAGCTTCTTCCCTCTAGCGAA
TGTGATCTCTGCCCACAAGCACTGGGTGAGGGCCTTGCCAGGACTTTTGTGCTTAGTATCTCTAAGTACATTTTC
TGTATTCTAGCAATTAGTGCCCTTATCTAACTACCCAGCTGCCTACCAATACTGGAACCTTCTTAAGGGTGGTGC
TCTCTTCTCTTGTCTGTTTGGCCCGTGGTGAGTAACGTGGAGCAGGTGCTGTGCTCAGTCAATATTGATGAAT
GAATGTGTACATGAGAGAACCAGAGTTGGTTCCTGGATAAATGGGCACATAAAAGCACAAAGTCCTTGG

902/6881
FIGURE 839

AAGGACTCCATGAAAGATGACAGAAGAAGTTATTGTGATAGCCAAGTGGGACTACACCGCCCAGCAGGACCAGGA
GCTGGACATCAAGAAGAACGAGCGGCTGTGGTTGCTGGACGACTCCAAGACGTGGTGGCGGGTGAGGAACGCGGC
CAACAGGACGGGCTATGTACCGTCCAACCTACGTGGAGCGGAAGAACAGCCTGAAGAAGGGCTCCCTCGTGAAGAA
CCTGAAGGACACACTAGGCCTCGGCAAGACGCGCAGGAAGACCAGCGCGCGGGATGCGTCCCCACGCCCAGCAC
GGACGCCGAGTACCCCGCCAATGGCAGCGGCGCCGACCGCATCTACGACCTCAACATCCCGGCCTTCGTCAAGTT
CGCCTATGTGGCCGAGCGGGAGGATGAGTTGTCCCTGGTGAAGGGGTGCGCGCTCACCGTCATGGAGAAGTGCAG
CGACGGTTGGTGGCGGGGAGCTACAACGGGCAGATCGGCTGGTTCCCTCCAACCTACGTCTTGAGGAGGTGGA
CGAGGCGGCTGCGGAGTCCCCAAGCTTCCTGAGCCTGCGCAAGGGCGCCTCGCTGAGCAATGGCCAGGGCTCCCG
CGTGCTGCATGTGGTCCAGACGCTGTACCCCTTCAGCTCAGTCACCGAGGAGGAGCTCAACTTCGAGAAGGGGGA
GACCATGGAGGTGATTGAGAAGCCGGAACGACCCCGAGTGGTGGAAATGCAAAAATGCCCGGGGCCAGGTGGG
CCTCGTCCCCAAAACCTACGTGGTGGTCCCTCAGTGACGGGCCTGCCCTGCACCCTGCGCACGCCCCACAGATAAG
CTACACCGGGCCCTCGTCCAGCGGGCGCTTCGCGGGCAGAGAGTGGTACTACGGGAACGTGACGCGGCACCAGGC
CGAGTGCGCCCTCAACGAGCGGGGCGTGGAGGGCGACTTCCTCATTAGGGACAGCGAGTCCTCGCCCAGCGACTT
CTCCGTGTCCCTTAAAGCGTCAGGGAAGAACAACACTTCAAGGTGCAGCTCGTGGACAATGTCTACTGCATTGG
GCAGCGGCGCTTCCACACCATGGACGAGCTGGTGGAACTACAAAAGGCGCCCATCTTCACCAGCGAGCACGG
GGAGAAGCTCTACCTCGTCAGGGCCCTGCAGTGACGGCGCCCGGCCCCACACTCGCCTCCCGGGCCCCACGGTG
GAGCTGCCCGCCCGGCCTGTGGCAGAGGCTCCTCCCGCGGGACGGCCCCGACGGCTTCTCTG

903/6881

FIGURE 840

MTEEVIVIAKWDYTAQQDQELDIKKNERLWLLDDSKTWVRVNAANRTIGYVPSNYVERKNSLKKGSLVKNLKDITL
GLGKTRRKSARDASPTPSTDAEYPANGSGADRIYDLNIPAFVKFAYVAEREDELSLVKGSRTVMKECSDGWWR
GSYNGQIGWFPSNYVLEEVDAAAESPFLSLRKASLSNGQGSRLHVVTLYPFSSVTEEEELNFEKGETMEVI
EKPENDPEWWKCKNARGQVGLVPKNYVVVLSGDPALHPAHAPQISYTGPFSSGRFAGREWYYGNVTRHQAECALN
ERGVGDFLIRDSESSPSDFSVSLKASGKNKHFKVQLVDNVYCIGQRRFHTMDELVEHYKKAPIFTSEHGEKLYL
VRALQ

904/6881
FIGURE 841

TGGAACCATGCTGCTTGAGGCTCTGGACTGCATCCTACCACCAACTCGTCCAAGTACAAGCCCTTGCACCTGCC
TCTCCAGGATGTCTACAAAATTGGTGGTATTGGTAGTGTTCTGTTGGCCGAGTGGAGACTGGTGTCTCAAACC
TGGTATGGTGGTCACCTTTGCTCCAGTCAACGTTACAACAGAAGTAAATCTGTCGAAATGCACCATGAAGCTTT
GAGTGAAGCTCTTCCTGGGGACAATGIGGGCTTCAATGTCAAGAATGTGTCTGTCAAGGATGTTTCGTCTGGCAA
CGTTGCTGGTGACAGCAAAAATGACCCACCAATGGAAGCAGCTGGTTTCACTGCTCAGGTGATTATCCTGAACCA
TCCAGGCCAAATAAGCACTGGCTATGCCCTGTATTGGATTGCCACACGGCTCATATTGCATGCAAGTTTGCTGA
GCTGAAGGAAAAGATTGATCACCGTTCTGGTAAAAAGCTGGAAGATGGCCCTAAATTCTTCTTGGGTTGCTTTG
CTGTTTCGTGATATGAGACAGATAGTT

905/6881

FIGURE 842

AAAAGCCAAAATGGGAAAGGAAAAGACTCATACCAACATTGTCGTCATTGGACACATAGATTTCGGGCAAGTCCAC
CACTACTGGCCATCTGATCTACAAAGGCGGTGGCATCGACAAAAGAACCATTGAAAAATTTGAGAAGGAGGCTGC
TGAGATGGGAAAGGGCTCCTTCAAGTATGCCTGGGTCTTGGATAAACTGAAAGCTGAGTGTGAACATGGTATCAC
CATTGATACCTCCTTGTGGAAATTTGAGACCAGCAAGTACTATGTGACTATCGTTGATGCCCCAGGACACAGAGA
CTTCATCAAAAACATGATTACAGAGACATCTCAGGCTGACTGTGCTGTCTGATTGTTGCTGCTGGTGTGGTGA
ATTTGAAGCTGGTATCTCCAAGAATGGGCAAACCCAAGAGCATGCCCTTCTGGCTTACACACTGGGTGTGAAACA
ACTAATTGTTGGTGTTAACAAAATGGATTCTACTGAGCCACCCTACAGCCAGAAGAGATATGAGGAAACTACAAA
ATTGGTGGTATTGGTAGTGTTCCTGTTGGCCGAGTGGAGACTGGTGTCTCAAACCTGGTATGGTGGTCACCTTT
GCTCCAGTCAACGTTACAACAGAAGTAAATCTGTGAAATGCACCATGAAGCTTTGAGTGAAGCTCTTCCTGGG
GACAATGTGG

906/6881

FIGURE 843

AACAGATCTTCATGAATGTCGCTATCTTTGAGGATGAAGTTTTTGCTGGAGTTACCACACACCAGGAACCTCTTTC
CACACAGCCTGCTGAGTGTGATTGCCAACTTCATCCCTTTCTCTGATCTCAACCAGAGTCCACGGAACATGTACC
AATGCCAGATGGGTAAGCAAACATATGGGCTTTCCACTTCTCACTTATCAAGACCGATCGGATAACAACTGTATC
GTCTTCAGACTCCTCAGAGTCCCTTGGTGAGACCTCCATGTATGATTATTATGACATGGATAACTATCCAATTG
GGACCAATGCCATCGTTGCTGTGATTCTTACACTGGCTATGATATGGAAGATGCCATGATTGTGAATAAGGCCT
CTTGGGAACGAGGCTTTGCCCATGGAAGTGTCTACAAGTCTGAGTTCATAGACCTCTCTGAAAAAATTAAACAAG
GAGATAGTAGCCTGGTGTGTTGGCATCAAACCTGGTGACCCACGCGTTCTGCAGAAGTTAGATGACGATGGATTGC
CGTTTATAGGAGCAAACTGCAGTACGGAGATCCGTATTACAGCTACCTCAACCTCAACACCGGGGAAAGTTTTG
TGATGTACTATAAGAGTAAAGAAAATTGTGTTGTGGATAACATCAAAGTGTGCAGTAATGACACTGGGAGTGGA
AATTCAAGTGTGTTTGCATCACTATGAGAGTGCCTCGGAACCCAACTATCGGAGATAAATTTGCCAGTCGCCATG
GGCAGAAGGGCATTTTAAGCAGATTGTGGCCGGCTGAGGACATGCCTTTTACTGAGAGTGGGATGGTCCAGACA
TTCTGTTCAATCCCCATGGTTTTCCATCCCGCATGACCATTGGGATGTTAATTGAGAGTATGGCCGGGAAGTCTG
CAGCTTTGCATGGTCTCTGCCATGATGCTACACCCTTCATCTTCTCAGAGGAGAACTCGGCCTTAGAATACTTTG
GTGAGATGTTAAAGGCTGCTGGCTACAATTTCTATGGCACCGAGAGGTTATATAGTGGCATCAGTGGGCTAGAAC
TGGAAGCAGACATCTTCATAGGAGTGGTTTATTATCAGCGCTTACGCCATATGGTCTCAGACAAAATTTCAAGTAA
GGACAACCTGGAGCCCGAGACAGAGTCACCAACCAGCCTATTGGGGGAAGAAATGTCCAGGGTGGAATCCGTTTTG
GGGAGATGGAACGGGATGCGCTTTTAGCTCATGGTACATCTTTTCTCCTTCATGACCGCTCTTCAACTGCTCAG
ATCGGTCGGTAGCCCATGTGTGTGAAGTGTGGCAGTTTACTCTCTCCACTGTTGGAGAAGCCACCCCTTCTTGG
TCTGCCATGCGCAACAGAAAATACAACGTACTCTGTGTAGTCTGCGAGTGACACTATCGATACTGTTTCTGTGCCT
TATGTTTTTTCGGTATTTTGTAGCTGAACGGCAGCTATGAACATCAAAGTGAAACTGGATGTTGTTTAACTTGAT
GTTGACCTTTTGGATTAAAGGGGACTATCAGATTAAAGCAAAATGTAATTTTAAATCAATGAAGATATCATTACCA
GGTTACTCTTGAGATTTTCAACGGTGTAGAAGTCTCAACCAAGACCTGAAAACCAAGTATGCAAGGTTTCTGA
ATCTCTCTGGTAGATTAACTATTGACAATGATTTTCTGTTATCTTTGTTCAAAAAGTTCATGTCTTCTCAAAATA
TGAAATATTGATAAATGGAAGAGCATACGGTGACAAGTCTCCTTTCCAACCCAGGTTCCCTACACCCTGCTCTC
AGCAGGCAGTGAGTGTACACACCTGTTAATCCATCTTGAGCAGGACAGTACTATACAAATAGAATGCAAGCTGT
AATGTAATTTTATATTTTCTTATAGCCACGTTGAAGTAAAAACAAACAGGTACAGTGTTTTTTACCAGCTTTATA
GAAGTACAGTTGTTACATATTTAATGAATACAATTTGATGGGTCTGACTATATGCACACACCTTTGATACCATCA
CCACAATCAGGGTAATAAACATACCTGTCTCCACAAGTTTCTCCTGCCCCCTTGTTTTTTGCTTTTGGTT
GCTGTTGAGTTTTTGTGTTTCTCTGTGGTAAGAACACTTAACTCAAGACCTACCCTCTTAACAAATCTTTAAG
TGCACGATATAGTATTGTTAATTCAGGCACCATGTTGTACAACAGATCTTTAGACCTTACTTGTCTGCATAAC
TGAAGCTTTATACCTGTTGAACAACTCTCCATTTCCCTGGCCCCCTAGCAACCACCTTCTACCCTGTTTCTATGA
GTTTGACTATTACAGATATCTCATATAGTGGGATCATGCAATATTTGTCCTGTGACTGGCTTATTTCACTTAGCA
TAGTGAATAAGATTATCCATTTTGAAGCCAGGCATGGTGTGTGCATCTATAGTCCCTGCTATTTGAGAGGC
TGAGGTGGGAGGATCATTTGAGTGCAGGAGTTCAAGGACAGCCTGGGTAATATAGGAAGACCCTGTCTTGAAGAC
CCTGACCTCAAGTGATCCACCCACCTCGGCCTCCGAAAGTGCTAGGATTACAGGTGTGAGCCACTGTGCCTGGCC
TCCGGTGAGTATTTTATATTTAGTCTACACTTCCATACTTGGCTTTTTTCTGCTTTTATGTTGATCTGCTTTTAT
AGCAGTGTGTAGAGTGCCACTTATGTTTTCTTCTGTGTACAGTATTTTATTGTATGGATTTACCATCCCTGT
GTATTTAAGTTGTTCCATTCTTTGGCCATTATAACTTTTTCTGCAAATATTCTGGTGACTTATCTTTGGCCATT
ATAAAGTGTGATAATAAAAAAAAAAAAAAAAAA

907/6881

FIGURE 844

MNVAIFEDEVFAGVTTHQELFPHSLLSVIANFIPFSDLNQSPRNMYQCQMGKQTMGFPLLTYYQDRSDNKLYRLQT
PQSPLVRPSMYDYYDMDNYPIGTNAIVAVISYTGDMEDAMIVNKASWERGFAHGVSVKSEFIDLSEKIKQGDSS
LVFGIKPGDPRVLQKLDDGLPFFIGAKLQYGDPIYSYLNLTGESFVMIYKSKENCVVDNIKVCSNDTGSQKFKC
VCITMRVPRNPTIGDKFASRHGQKGILSRLWPAEDMPFTESGMVPDILFNPHGFPSRMTIGMLIESMAGKSAALH
GLCHDATPFIIFSEENSALEYFGEMLKAAGYNFYGTERLYSGISGLELEADIFIGVVYYQRLRHMVSDKFQVRTTG
ARDRVNTQPIGGRNVQGGIRFGEMERDALLAHGTSFLLHDLRFNCSDRSVAHVCEVWQFTLSTVGEATPFLVCHA
QQKIQLYSV

908/6881
FIGURE 845

GAGCTGTCCCCGGTGCCGCCGACCCGGGCCGTGTGCCCGTGGCTCCAGCCGCTGTGCGCTCGATCTCCTCGTCTC
CCGCTCCGCCCTCCCTTTTCCCTGGATGAACCTGCGTCCTTTCTTTCTCCGCCATGGAATTCGTCTCCGTGCTT
TTAGCCCTCCTGAGCCAAAGAAACCCAGACAACAGATGCCCATACGCAGCGTATAGCAGTAACCTCCCAAGCTCG
GTTTCTGTGCCGTAGTTTACAGTATTTAATTTTATATAATATATATTATTTATATAGCATTTTTGATACCTCAT
ATTCTGTTTACACATCTTGAAAGGCGCTCAGTAGTTCTCTTACTAAACAACCACTACTCCAGAGAATGCAACGC
TGATTACCAGTACTACAGCTGCTACCGCCGCTTCTGGTCTTTGGTGGACTACCTATGGATGCTCATCTGGGCT
TCATTATTGCATTTGTCTTGGCATTCTCCGTGGGAGCCAATGATGTAGCAAATTCCTTTGGTACAGCTGTGGGCT
CAGGTGTAGTGACCCTGAAGCAAGCCTGCATCCTAGCTAGCATCTTTGAAACAGTGGGCTCTGTCTTACTGGGGG
CCAAAGTGAGCGAAACCATCCGAAGGGCTTGATTGACGTGGAGATGTACAACCTCGACTCAAGGGCTGCTGATGG
CCGGCTCAGTCAGTGCTATGTTTGGTCTGTCTGTGTGGCAACTCGTGGCTTCGTTTTGAAGCTCCCTATTTCTG
GAACCCATTGTATTGTTGGTGCAACTATTGGTTTCTCCCTCGTGGCAAAGGGGCAGGAGGGTGTCAAGTGGTCTG
AACTGATAAAAAATTGTGATGCTTGGTTCTGTGCCCCACTGCTTCTGGAATTATGCTGGAATTTTATCTTCC
TGGTTCGTGCATTTCCTCCATAAGGCAGATCCAGTTCCTAATGGTTTGGCAGCTTTGCCAGTTTTCTATGCCT
GCACAGTTGGAATAAACCTCTTTCCATCATGTATACTGGAGCACCGTTGCTGGGCTTTGACAAACTTCCTCTGT
GGGGTACCATCCTCATCTCGGTGGGATGTGCAGTTTCTGTGCCCTTATCGTCTGGTTCCTTTGTATGTCCAGGA
TGAAGAGAAAAATTGAACGAGAAATAAAGTGTAGTCCTTCTGAAAGCCCCCTAATGAAAAAAGAATAGCTTGA
AAGAAGACCATGAAGAAACAAAGTTGTCTGTTGGTGATATTGAAAACAAGCATCCTGTTTCTGAGGTAGGGCTG
CCACTGTGCCCTCCAGGCTGTGGTGGAGGAGAGAACAGTCTCATTCAAACTTGGAGATTTGGAGGAAGCTCCAG
AGAGAGAGAGGCTTCCAGCGTGGACTTGAAAGAGGAAACCAGCATAGATAGCACCGTGAATGGTGCAGTGCAGT
TGCCTAATGGGAACCTTGTCCAGTTCAGTCAAGCCGTGAGCAACCAAATAAACTCCAGTGGCCACTACCAGTATC
ACACCGTGCATAAGGATTCCGGCCTGTACAAAGAGCTACTCCATAAATTACATCTTGCCAAGGTGGGAGATTGCA
TGGGAGACTCCGGTGACAAACCTTAAGGCGCAATAATAGCTATACTTCCTATACCATGGCAATATGTGGCATGC
CTCTGGATTCAATCCGTGCCAAAGAAGGTGAACAGAAGGGCGAAGAAATGGAGAAGCTGACATGGCCTAATGCAG
ACTCCAAGAAGCGAATTCGAATGGACAGTTACACCAGTTACTGCAATGCTGTGTCTGACCTTCACTCAGCATCTG
AGATAGACATGAGTGTCAAGGCAGAGATGGGTCTAGGTGACAGAAAAGGAAGTAATGGCTCTCTAGAAGAATGGT
ATGACCAGGATAAGCCTGAAGTCTCTCTCCTCTTCCAGTTCCTGCAGATCCTTACAGCCTGCTTTGGGTCAATCG
CCCATGGTGGCAATGACGTAAGCAATGCCATTGGGCCTCTGGTTGCTTTATATTTGGTTTATGACACAGGAGATG
TTTCTTCAAAAGTGGCAACACCAATATGGCTTCTACTCTATGGTGGTGTGGTATCTGTGTTGGTCTGTGGGTTT
GGGGAAGAAGAGTTATCCAGACCATGGGGAAGGATCTGACACCGATCACACCCTCTAGTGGCTTCAGTATTGAAC
TGGCATCTGCCCTCACTGTGGTGATTGCATCAAATATTGGCCTTCCCATCAGTACAACACATTGTAAAGTGGGCT
CTGTTGTGTCTGTTGGCTGGCTCCGGTCCAAGAAGGCTGTTGACTGGCGTCTCTTTCGTAACATTTTATGGCCT
GGTTTGTACAGTCCCCATTTCTGGAGTTATCAGTGCTGCCATCATGGCAATCTTCAGATATGTCATCCTCAGAA
TGTAAGCTGTTTGGAGATTAAATTTGTGTCAATGTTTGGGACCATCTTAGGTATTCCTGCTCCCCTGAAGAATG
ATTACAGTGTAAACAGAAGACTGACAAGAGTCTTTTATTTGGGAGCCAGAGGAGGAAGTGTACTTGTGCTAT
AACTGCTTTTGTGCTAAATATGAATTGTCTCAAATTAGCTGTGTAAATAGCCCGGGTTCCACTGGCTCCTGCT
GAGGTCCCCTTTCTTCTGGGCTGTGAATTCCTGTACATATTTCTCTACTTTTGTATCAGGCTTCAATTCATT
ATGTTTTAATGTTGCTCTGAAGATGACTTGTGATTTTTTTTTCTTTTTTAAACCATGAAGAGCCGTTTGACA
GAGCATGCTCTGCGTTGTTGGTTTACCAGCTTCTGCCCTCACATGCACAGGGATTTAACAACAAAAATATAACT
ACAACCTCCCTTGTAGTCTCTTATATAAGTAGAGTCTTGGTACTCTGCCCTCCTGTCTAGTGTGCGAGATCTA
TTGGCATATTCCGGAGCTTCTTAGAGGGATGAGGTTCTTTGAACACAGTGAAAATTTAAATTAGTAACCTTTTTG
CAAGCAGTTTATTGACTGTTATTGCTAAGAAGAAGTAAGAAAGAAAAGCCTGTTGGCAATCTTGGTTATTTCTT
TAAGATTTCTGGCAGTGTGGGATGGATGAATGAAGTGAATGTGAACTTGGGCAAGTTAAGTGGGACAGCCTTC
CATGTTCAATTTGTCTACCTCTTAACTGAATAAAAAAGCCTACAGTTTTTAG

909/6881

FIGURE 846

MATLITSTTAATAASGPLVDYLWMLILGFIIAFVLAFSVGANDVANSFGTAVGSGVVILKQACILASIFETVGSV
LLGAKVSETIRKGLIDVEMYNSTQGLLMAGSVSAMFGSAVWQLVASFLKLPISGTHCIVGATIGFSLVAKGQEGV
KWSelikivMSWfVSPllSGIMSGILFFLVRAFI LHKADPVPNGLRALPVFYACTVGINLFSIMYTGAPLLGFDK
LPLWGTILISVGCavFCALIVWFFVCPRMKRKIEREIKCSPSESPLMEKKNSLKEDHEETKLSVGDIENKHPVSE
VGPA TVPLQAVVEERTVSFKLGdLEEAPERERLPSVDLKEETSIDSTVNGAVQLPngNLVQFSQAVSNQINSSGH
YQYHTVHKDSGLYKELLHKLHLAKVGDCMGDSGDKPLRRNNSYTSYMAICGMPLDSFRAKEGEQKGEEMEKLW
PNADSKKRIRMDSYTSYCNVSDLHSASEIDMSVKAEMGLGDRKGSNGSLEEWYDQDKPEVSLLFQFLQILTACF
GSFAHGNDVSNAIGPLVALYLVDYTDGVSSKVATPIWLLLYGGVGICVGLWVWGRRVIQTMGKDLTPITPSSGF
SIELASALTTVIASNIGLPIS TTHCKVGSVSVSGWLRSKKAVDWRLFRNIFMAWFVTVPISGVISAAIMAIFRYV
ILRM

910/6881
FIGURE 847

ACCAACCTCTTCGAGGCACAAGGCACAACAGGCTGCTCTGGGATTCTCTTCAGCCAATCTTCATTGCTCAAGTGT
CTGAAGCAGCCATGGCAGAAGTACCTGAGCTCGCCAGTGAAATGATGGCTTATTACAGTGGCAATGAGGATGACT
TGTTCTTTGAAGCTGATGGCCCTAAACAGATGAAGTGCTCCTTCCAGGACCTGGACCTCTGCCCTCTGGATGGCG
GCATCCAGCTACGAATCTCCGACCACCACTACAGCAAGGGCTTCAGGCAGGCCGCGTCAGTTGTTGTGGCCATGG
ACAAGCTGAGGAAGATGCTGGTTCCCTGCCCACAGACCTTCCAGGAGAATGACCTGAGCACCTTCTTTCCCTTCA
TCTTTGAAGAAGAACCTATCTTCTTCGACACATGGGATAACGAGGCTTATGTGCACGATGCACCTGTACGATCAC
TGAAGTGCACGCTCCGGGACTCACAGCAAAAAAGCTTGGTGATGTCTGGTCCATATGAACTGAAAGCTCTCCACC
TCCAGGGACAGGATATGGAGCAACAAGTGGTGTCTCCATGTCCTTTGTACAAGGAGAAGAAAGTAAAGCAAAA
TACCTGTGGCCTTGGGCCTCAAGGAAAAGAATCTGTACCTGTCCTGCGTGTGAAAGATGATAAGCCCACTCTAC
AGCTGGAGAGTGTAGATCCCAAAAATTACCCAAAGAAGAAGATGGAAAAGCGATTGTCTTCAACAAGATAGAAA
TCAATAACAAGCTGGAATTTGAGTCTGCCCAGTTCCCAACTGGTACATCAGCACCTCTCAAGCAGAAAAACATGC
CCGTCTTCTGGGAGGGACCAAGGCGGCCAGGATATAACTGACTTCACCATGCAATTTGTGTCTTCTTAAAGAG
AGCTGTACCCAGAGAGTCTGTGCTGAATGTGGACTCAATCCCTAGGGCTGGCAGAAAGGGAACAGAAAGGTTTT
TGAGTACGGCTATAGCCTGGACTTTCTGTGTCTACACCAATGCCCAACTGCCTGCCTTAGGGTAGTGCTAAGA
GGATCTCCTGTCCATCAGCCAGGACAGTCAGCTCTCTCCTTTCAGGGCCAATCCCCAGCCCTTTTGTGAGCCAG
GCCTCTCTCACTCTCCTACTCACTTAAAGCCCGCTGACAGAAACCACGGCCACATTTGGTTCTAAGAAACCTT
CTGTCAATTCGCTCCACATTCTGATGAGCAACCGCTTCCCTATTTATTTATTTATTTGTTTGTGTTTATTCA
TTGGTCTAATTTATTCAAAGGGGGCAAGAAGTAGCAGTGTCTGTAAGAGAGCCTAGTTTTTAATAGCTATGGAAT
CAATTCAATTTGGACTGGTGTGCTCTCTTTAAATCAAGTCCTTAATTAAGACTGAAAATATATAAGCTCAGATT
ATTTAAATGGGAATATTTATAAATGAGCAAATATCATACTGTTCAATGGTTCTGAAATAAACTTCTCTGAAG

911/6881
FIGURE 848

MAEVP ELASEMMAYYS GNEDDLFFEADGPKQMKCSFQDL DLCPLDGGIQLRISDHHYSKGF RQAASVVVAMDKLR
KMLVPCPQTFQENDLSTFFPFIFEEPIFFDTWDNEAYVHDAPVRSINCTLRDSQQKSLVMSGPYELKALHLQGG
DMEQQVVF SMSFVQGEESNDKIPVALGLKEKNLYLSCVLKDDKPTLQLESVDPKNYPKKKMEKR FVFNKIEINN K
LEFESAQFPNWYIST SQAENMPVFLGGTKGGQDITDFTMQFVSS

912/6881
FIGURE 849

AGGCACCATGACTCCTGTGAGGATGCAGCACTCCCTGGCAGGTCAGACCTATGCCGTGCCCTCATCCAGCCAGA
CCTGCGGTGAGAGGAGGCCGTCCAGCAGATGGCAGATGCCCTGCAGTACCTGCAGAAGGTCTCTGGAGACATCTT
CAGCAGTGAGAGCGGCCTTGCTGCGGCCAAAGAACATGGAAGCATGAGTGGGGTGATGTGTGCCTTAAAGCAT
CAGACACTTGGGCCTCGGGCATCAGGAGCCAGCCACAGGGATGTCTGGGGAAATGGCGTTCCATGAGATGCAAGC
ACACAAGAATGCACTTGGCACATCTGGGGAACAGCAGGCAGCTGATATCACTGGGCCCACCCTGCACCAGGGAGG
ATGGAAGCAGGTAGAGCAGAGCCGGAGCCAGGTGCAGGCCATTGGAGAGAAGGTCTCCTTGGCCCAGGCCAAGAT
TGAGAAGATCAAGGGCAGCAAGAAGGCCATCAAGGTGTTCTCCAGTGCCAAGTACCCTGCTCCAGAGCGCCTGCA
GGAATATGGCTCCATCTTCAGGGCGCCCAGGACCCTGGCCTGCAGAGACGCCCCCGCCACAGGATCCAGAGCAA
GCACCGCCCCCTGGACGAGCGGGCCTGCAGGAGAAGCTGAAGGACTTTCCTGTGTGCGTGAGCACCAGCCGGA
GCCCAGGACGATGCAGAAGAGGGACTTGGGGGTCTTCCCAGCAACATCAGCTCTGTCAGCTCCTTGCTGCTCTT
CAACACCACCTAGAACCT

913/6881
FIGURE 850

AGCGCGGGGAACCGGAGGCTGCAGGATGGTCAAGCTGACGGCGGAGCTGATCGAGCAGGCGGCGCAGTACACCA
ACGCAGTGC GCGACCGGGAGCTGGACCTCCGGGGGTATAAAATTCCCGTCATTGAAAATCTAGGTGCTACGTTAG
ACCAGTTGATGCTATTGATTTTTCTGACAATGAGATCAGGAACTGGATGGTTTTCTTTGTGAGAAGACTGA
AAACATTGTTAGGGAACAACAACAGAATATGCCGTATAGGTGAGGGACTTGATCAGGCTCTGCCCTGTCTGACAG
AACTCATTCTACCAATAATAGGCTCGTGGAACTGGGTGATCTGGACCCTCTGGCATCTCTCAAATCGCTGACTT
ACCTAAGTATCCTAAGAAATCTGGTAACCAATAAGAAGCATTACAGATTGTATGTGATTTATAAAGTTCCGCAAG
TCAGAGTACTGGATTTCCAGAAAGTAAACTAAAAGTTTTAATCCAGGTGCTGGTTTGCCAACTGACAAAAAGAA
AGGTGGGCCATCGCCAGGGGATGTAGAAACAATCAAGAATGCTATAGCAAATGCTTCAACTCTGGCTGAAGTGGA
GCGGCTGAAGGGGTTGCTGCAGTCTGGTCAGATCCCTGGCAGAGAACGCAAATCGGGGCC

914/6881
FIGURE 851A

GATGGGCTGTGCAAAGGGCGCGGCTGCTGAGGACAGGGCTGTGGCTGGAGAACCTGCTGCTTCCTGAAGCCGAGC
CCCACCCTAAAGGACCGAGAGCACAGGCCATGGGAGAGGAATCGAGTAGGGATGCAGAGCCTGGTGGCACTGTCC
TGAGAGAGGAGGAGGCTGGGCAGGGCCCCGGCGCTGGAGAGGTGCCCTGCATATGGGACCTCTCAGAGGTGCACA
AGAGCTGTGTGCCAGGTACCCCTGTCTCCTCTCCTCCTCCAGCCCCAGCCTACTCTCCTAATGGTCTGTGTCT
TTCAGGGATGAGATTTACTGCCAGATCTGCAAGCAGCTCTCGGAGAACTTCAAACAGCAGCCTGGCCCCGGGC
TGGATCCTGCTCAGCCTCTGCCTCGGCTGCTTCCACCCTCAGAGAGGTTTCATGAAGTATCTACTGAACTTCATC
GGCCAAAGGGCCGGCAGCTACGGCCCCCTTCTGTGCCGAGCGCCTGAGACGCACCTATGCCAATGGGGTGCCTGCG
GAGCCCCCACCCTGGCTGGAGCTGCAGGCTGTCAAGTCCAAGAAGCACATCCCCATCCAAGTCATCTTGGCCACT
GGAGAGAGCCTAACCGTCCCCGTGGACTCAGCCTCCACATCTCGGGAAATGTGCATGCACATCGCTCACAAGCAG
GGCCTCAGCGACCACCTGGGCTTCTCCCTCCAGGTGCGCGTGTACGACAAGTTCTGGTCCCTGGGCAGCGGGCGC
GACCACATGATGGATGCCATCGCCCGGTGTGAGCAGATGGCCAGGAGAGGGGCGAGAGCCAGCGCCAGTCACCC
TGGCGCATCTACTTCCGGAAGGAATTTCTTACCCCTGGCAGCACTTCCCGGAGGACCTGTCTCAGCACCAGCTT
ATTTACCGCCAGTCTCCGAGGAGTCTGGTCTGGCGAGTACAGCTTCGAGAAGGAGGAAGAGCTGGTTGAGCTG
CTGGCCCCGCACTGCTACGTGCAGCTCGGCGCCTCAGCAGAGAGCAAGGCTGTCCAGGAGCTGCTGCCCAGCTGC
ATCCCCCACAAGCTGTACAGGACCAAGCCCCAGACAGGTGGGCGAGCCTCGTCACTGCCGCTGCGCCAAGGCC
CCATACACTCAGAAGCAAGTCACACCCTGGCCGTGCGAGAGCAGGTGGTGGACGCCGCCCGCTGCAGTGGCCG
CTGCTCTTCTCCGGCTCTTTCGAAGTCAACACTCTCAGGCCCCCCGCTGCCAAGACGCAGCTGATCTTGGCT
GTTAACTGGAAGGGGCTTTGCTTCTTGGACCAGCAGGAGAAGATGCTGCTGGAACCTCTCTTCCAGAGGTCATG
GGTCTGGCCACCAACAGGGAGGGCCAGGGCGGGCAGAGGCTGCTGCTCTCCACGATGCATGAGGAGTACGAGTTT
GTGTACCCAGCAGTGTGGCCATCGCTGAGCTGGTGGCCCTGTTCCTGGAGGGCCTGAAGGAGAGGTCCATTTTC
GCCATGGCCCTGCAGGACAGGAAGGCCACAGATGACACCACCTCCTGGCCTTCAAGAAGGGGGACCTGTTGGTC
CTCACAAGAAGCAGGGGCTGCTGGCCTCTGAGAACTGGACCCTCGGCCAGAACGACAGGACAGGCAAGACGGGG
CTGGTGGCCATGGCCTGCCTCTACCCATCCCCACGGTCACTAAGCCCTCGGCACAGCTGCTGAGCTTGCTTGGC
ATGTCACCAGAGAAGAGGAAGCTGGCGGCTCAGGAGGGGCAAGTTCACAGAGCCACGTCCTGAGGAGCCACCCAAG
GAAAAGCTGCACACCCTGGAGGAGTTCTCCTATGAGTTCTTCAGGGCTCCAGAGAAGGACATGGTGAGCATGGCC
GTGCTGCCCCCTGGCCCGTGGCCGCTGGCCACCTGTGGGCTATTCTCGCAGCCGCTGCGACAGCCGCTGCTCAAG
CGAGTCCACGCCAACGTGCACCTCTGGGACATCGCTGCCAGATCTTTGTCGCCATCTCCGGTACATGGGCGAC
TACCCTTCTCGGCAGGCTGGCCACCCTGGAGCTCACCGACCAGATCTTCACACTGGCCCTGCAGCACCCTGGCC
CTCCAGGACGAGGTCTACTGCCAGATCCTGAAGCAGCTGACGCACAACCTCAACAGGCACAGCGAAGAGCGGGG
TGGCAGCTGCTGTGGCTGTGCACGGGCTCTTCCCGCCAGCAAGGGGCTGCTGCCCCATGCCAGAAGTTTATA
GACACTCGGAGGGGGAAGCTGCTGGCCCCGACTGCAGCGCCGAATCCAGAAGGTCTGAGGACGGGGCCCCGG
AAGCAGCCCCCGCACCAAGGTGGAGGTGGAGGCGCAGAGCAGAACGTCTCCGCATCTGCCACAAGATCTACTTC
CCCAATGACACCAGTGAGATGCTGGAGGTGGTTGCCAACACACGGGTGCGGGATGTGTGTGACAGCATTGCCACC
AGGCTGCAGCTGGCCTCCTGGGAGGGCTGCAGCCTCTCATCAAGATTTAGACAAGGTGGGCCGGGCTGGGGCT
GGGCAGACGGTGGGCGGACGGGCAAGTGGAGCGAGGCCCTAGGTGCTGCTGTGGGGGCTCAGCCTACCAGGGGCA
CCCATGCTGGACCAGGCTGCACGGCCAGGGCTGCTGGGACAGAGGTAGGTGGGGTCTTGAGCAGAGCCAGCTCC
ACACGCCAATAGCCACGCACAGCCGACTCCAGCGCAGCACCAGGCCGTAGGCGGCCACTGGACCAGAACCAG
CAGGTTCTCAGGAGGACAGTCCCCGGAAGCCACCAACTTCCCTGTACCTTCCCTTCCCAAGTTCATCAGCCAG
AAGGAGGGGAGACTTCTTCTTTGATTCTTGAGGGAGGTGTCTGACTGGGTGAAGAAGAACAAGCCCCAGAAAGAA
GGGGCCCCCGTGACGCTCCCCCTACCAGGTGTACTTCATGCGGAAATTGTGGCTCAACATATCTCCAGGGAAGGAT
GTGAATGCAGACACCATACTCCATTACCACCAGGAGCTGCCCAAGTACCTGCGCGGATTCCACAAGTGTTCGCGG
GAGGATGCCATCCACCTGGCGGGCTCATCTACAAGGCCAGTTCAACAACGACCGGTCCAGCTGGCTAGTGTG
CCCAAGATCCTGAGGGAAGTGGTGCCTGAGAACCTCACACGCTGATGTCTCGGAGGAGTGGAAAAAGAGCATC
CTTCTAGCCTATGACAAGCATAAGGACAAGACAGTGGAGGAGGCCAAGGTGGCCTTCTGAAGTGGATCTGCCGG
TGGCCACCTTCGGATCCGCCTTCTTCGAGGTGAAGCAAACCTCGGAGCCTTCTTACCCGACGTCATCTCATC
GCCATCAACCGACATGGGGTTCTGCTCATCCACCCCAAGACCAAGGACCTGCTCACCACCTATCCCTTACCAAG
ATCTCCAGCTGGAGCAGCGGCAGCACCTACTTCCACATGGCGCTGGGAGCCTGGGCGGTGGCAGCCGCTGCTG
TGCGAGACCTCCCTGGGCTATAAGATGGATGACCTGCTGACCTCATATGTGCAGCAGCTCCTGAGTGCCATGAAC

915/6881
FIGURE 851B

AAGCAGCGGGGCTCCAAGGCCCCAGCCCTGGCCAGCACCTAGCAGCGGATGCTGGCGTGTCTGCTCAGGCGCCCT
TCCCGACCTCTAGCCTGGCGGCACCTTCCCAGGCCCTCTCAACCCAGGGCCTGTCCTTGGCGGGCAGCCTTCCAT
GCTGCCCCCATAAAAGCCCACTCAGCCCCGCAGGCGGCCCCCTCTGTCCTGGGCGCTGCCAGGGAGGCCAAA
AGACGGGCCCAGAATGGGGTCGGGAGTCTCGGACCCCCAGGCTATTGGTGGATGACTGACTGACAGGACACCTCC
CAACCCACCCACCCACCAGAATGTTCAATAAAAACCTCCTGGAGCAGGA

916/6881
FIGURE 852

MKYLLNFIGQGPATYGPFCERLRRTYANGVRAEPPTWLELQAVKSKKHIPIQVILATGESLTPVDSASTSREM
CMHIAHKQGLSDHLGFSLQVAVYDKFWSLGSRDHMDAIARCEQMAQERGESQRQSPWRIYFRKEFFTPWHDSR
EDPVSTELIYRQVLRGVWSGEYSFEKEEELVELLARHCYVQLGASAESKAVQELLPSCHIPKLYRTKPPDRWASL
VTAACAKAPYTQKQVTPLAVREQVVDAAARLQWPLLF SRLFEVITLSGPRLPKTQLILAVNWKGLCFDQOEKMLL
ELSFPEVMGLATNREAQGGQRLLLSTMHEEYEFVSPSSVAIAELVALFLEGLKERSIFAMALQDRKATDDTTLLA
FKKGDLLVLTKKQGLLASENWTLGQNDRTGKTGLVPMACLYTIPTVTKPSAQLLSLLAMSPEKRKLAQEGQFTE
PRPEEPPKEKLHTLEEFSEYEFFRAPEKDMVSMVLP LARARGHLWAY SCEPLRQPLLKRVHANVDLWDIACQIFV
AILRYMGDYP SRQAWPTLELTDQIFTLALQHPALQDEVYQCILKQLTHNSNRHSEERGWQLLWLCTGLFPFSKGL
LPHAQKFIDTRRGKLLAPDCSRRIQKVLRTGPRKQPPHQVEVEAAEQNVSRICHKIYFPNDTSEMLEVVANTRVR
DVCDSIATRLQLASWEGCSLFIKISDKVGRAGAGQTVGGRAVSEALGAACGGLSLPGAPMLDQAARPGLLGQR

917/6881
FIGURE 853

AAGTGCATGACTCAGGACTGGGTGGGAGGGGGCTGTCCCTGAGGCCCTGGTCTCCCCAGCCAAGTCTGGCAGG
GGGATGCTGTTCTTCAGGATCCCATGACCTGGGGCTCTTGGGGAGCCCCAGCCCGAGAGTGACCTGTGGCTTTG
CTGGGGAAAGGGGCTTGGAGGGTACAGGGCAGTGTCCAGGATTGCAGTCTCAAATCCGAGGCCCCACAGGGGAGCT
GCCTCTGTGATCGCCATTAAGGCTGTGGCCTGGGGTCTCAGGGCAGCCCAGGAACCAAAATAGCCTGTAACTTG
TCACTTTGTAAGTCTGCCCTCCAGACACACTTCCCTTTCTTCTCCCAAGCCTCTCTTCCCAGTTAGCTCCCT
GACTTGGAGTCACCTCTAAGCCTTGGCCCAGGCCCTCTGCAGTGAGATGGAGGGACTCATCCCCCGAGAGTGG
CATCTGCAGGCCCCACTGGCTGCCTGTGGTCACCCCTAGGGTTCTAGCGGGGCTGCCTCCATTTTCTCAGAGC
ATCCCTGCCAGCTGGCTGGTTCTACTCTCCTGTTTTCTAGGAAAAGAAAGTGGGGAGAGACAGGAACAGGGTCC
GGAGCAGCCCCAGCAGCAGCATTGAGGTGGAGCAATATGTCGGACGCCTTGGCCAACGCCGTGTGCCAGCGCTGC
CAGGCCCCGCTTCTCCCCCGCGAGCGCATTGTCAACAGCAATGGGGAGCTGTACCATGAGCACTGCTTCGTGTGT
GCCAGTGCTTCCGGCCCTTCCCGAGGGGCTCTTCTATGAGTTTGAAGCCGGAAGTACTGCGAACACGACTTC
CAAATGCTGTTTGTCCGTGCTGTGGATCCTGCGGTGAGTTTCATCATTGGCCGCGTCATCAAGGCCATGAACAAC
AACTGGCACCCGGGCTGCTTCCGCTGCGAGCTGTGTGATGTGGAGCTGGCTGACCTGGGCTTTGTGAAGAATGCC
GGCAGGCATCTCTGCCGGCCTTGCCACAACCGTGAGAAGGCCAAGGGCCTGGGCAAGTACATCTGCCAGCGGTGC
CACCTGGTCATCGACGAGCAGCCCCCTCATGTTTCAGGAGCGACGCCTACCACCTGACCACTTCAACTGCACCCAC
TGTGGGAAGGAGCTGACAGCCGAGGCCCGGAGCTGAAGGGTGAGCTCTACTGCCTGCCCTGCCATGACAAGATG
GGCGTCCCCATCTGCGGGGCTGCGCGCGGCCCATCGAGGGCCGAGTGGTCAACGCGCTGGGCAAGCAGTGGCAC
GTGGAGCACTTTGTCTGTGCCAAGTGTGAGAAGCCATTCTGGGGCACCAGCACTATGAGAAGAAGGGCCTGGCC
TACTCGGAGACTCACTACAACAGCTCTTCGGGGACGTCTGCTACAACCTGCAGCCATGTGATTGAAGGCGATGTG
GTGTGCGCCCTCAACAAGGCCTGGTGTGTGAGCTGCTTCTCCTGCTCCACCTGCAACAGCAAGCTCACCTGAAG
AACAAGTTTGTGGAGTTTCGACATGAAGCCCGTGTGTAAGAGGTGCTACGAGAAGTTCCCGCTGGAGCTGAAGAAG
CGGCTGAAGAAGCTGTGCGGAGCTGACCTCCCGCAAGCCAGCCCAAGGCCACAGACCTCAACTCTGCCTGAAGGC
CCTCTTGCGCAGTGCCCTCTCGGCCCCCTCCGCCTTCTCCCTCCTGCTGTCCATGCTTGGCCCCCTCGTCCCCATC
CACCTGTGCCCCCTCCGCATCTTACCCTCCCTTTCTCTTTCTCATTGCCTTCTCCCTTCCCTGTTCCTCATCTCTG
CCTTCCCCATGTCTCTCCTCTCCTTGGCCGTGGCTTCTGTCTGTGAGGAGGCAGGAGCTGGGGAGTGGGAGCCTA
TGACCCACAGTCTGACAGCCATGTCCACCTGTGCCACAGCTTCCGCCCACAGACCTCCAGGGACAGGAGCAAT
TGACCCACAGCTCCCCGCTGGCCTGGCCCTCCCCAGGCGGCTCAGTGGCTCATGCTGTCTGTGAGAGCCCCTG
CCCCAGAGCGGCCCCACTAAGCGCATGTGGCTCCTGGGCTACCCACAGCCAGGGCAGCCTGCTGGAGCCACAGGG
CCAGGGCCATGCAGATGGAGGCCTCTGGGAGCCACCTCCAATCCCTCACCCTCACTCAACCAGTGGCACAGTGT
CCTTGTGCCACACTGAGCCAGCAAGTCTGTGTCCACACCCACAAGCTACCTGGAGGGACAGGACCCACCTCC
ATCCTTCGGAAGGCCTTCTGGAATCCACCTTGGCCTCCGCCCTCGGTTCCGCCCGCCCCCTCTCCCCCGACC
TTGGGGCTTGTGTGAGCCCTTGGGTGGGGCCAGGAGGAGGTGATGGCGTCAGAGGAGGTGTGGTCAGAGGTGAC
TTGTTCCACCTCCAGGGAGGACGTTCTGTTCTCGGCCAGCGCAGACCTGGTGTGTTGTTGTTGGGCTCACGCTT
GCACAATGAAGGCTTGTTACAC

918/6881
FIGURE 854

CCGACACCCACGGGCGGAGATCACCTGCTGCCCCGCAGACCCCTGTCCCTTCCTCCCGGACCAGCAGCTAGAGGA
TGTCCAAACGGAGTTGGTGGGCTGGATCCAGAAAGCCCCAAGAGAGATGCTGAAACTCTCAGGCTCTGACTCCA
GCCAAAGCATGAATGGCCTTGAAGTGGCTCCCCCAGGTCTGATCACCAACTTCTCCCTGGCCACGGCAGAGCAAT
GTGGCCAGGAGACGCCACTGGAGAACATGCTGTTGCGCTCCTTCTACCTTCTGGATTTTATCCTGGCTTTAGTTG
GCAATACCCTGGCTCTGTGGCTTTTTCATCCGAGACCACAAGTCCGGGACCCCGGCCAACGTGTTCTGTATGCATC
TGGCCGTGGCCGACTTGTGCTGCGTGTGGTCTGCCCACCCGCTGGTCTACCACTTCTCTGGGAACCACTGGC
CATTTGGGGAAATCGCATGCCGTCTCACGGCTTCTCTTCTACCTCAACATGTACGCCAGCATCTACTTCTCTCA
CCTGCATCAGCGCCGACCGTTTCTGGCCATTGTGCACCCGGTCAAGTCCCTCAAGCTCCGCAGGCCCCCTCTACG
CACACCTGGCCTGTGCCTTCTGTGGGTGGTGGTGGCTGTGGCCATGGCCCCGCTGCTGGTGAGCCACAGACCG
TGCAGACCAACCACACGGTGGTCTGCCTGCAGCTGTACCGGAGAAGGCCTCCCACCATGCCCTGGTGTCCCTGG
CAGTGGCCTTACCTTCCCGTTTCATACCACGGTTCACCTGCTACCTGCTGATCATCCGCAGCCTGCGGCAGGGCC
TGGGTGTGGAGAAGCGCTCAAGACCAAGGCAGTGCGCATGATCGCCATAGTGTGGCCATCTTCTGGTCTGCT
TCGTGCCCTACCACGTCAACCGCTCCGTCTACGTGCTGCACTACCGCAGCCATGGGGCCTCCTGCGCCACCCAGC
GCATCCTGGCCCTGGCAAACCGCATCACCTCCTGCCTCACCAGCCTCAACGGGGCACTCGACCCCATCATGTATT
TCTTCGTGGCTGAGAAGTTCCGCCACGCCCTGTGCAACTTGCTCTGTGGCAAAAGGCTCAAGGGCCCGCCCCCA
GCTTCGAAGGGAAAACCAACGAGAGCTCGCTGAGTGCCAAGTCAGAGCTGTGAGCGGGGGGCGCCGTCCAGGCCG
AGCGCAGACTGTTTAGGACTCAGCAGACCCAGCAAGAGGCATCTGCCCTTTCCCCAGCCACCTCCCCAGCAAGCA
ACCTGAAATCTCAGCAGATGCCCCACATTTCTCTAGATCGCCTAGTCTCAACCCATAAAAAGGAAGAACTGACAA
AGGGGATCCATCGGCCACCCCTCTGCAGGGGCTTGTGATGGCTACAATGGCTCCTAGACACTCAACGACTTCATC
TGTGGCAGGGAGAGAGGAGGCCGGAAGAACAACCCCTGAACAATGGAGGCCTTTCTTTCCCGCTAGGCTCCCAGC
CTCCTTCCCGCTACAGAATCGCTCATCGGCAGGGCTCAGCAGAAAGACCCTGAAGGCAGGCTGCAAATGACCCAG
AAGAGGGACCTGGGAGTCTTGGTGGGGACGGGGAGGGAGTCTCAATACTCCTTTGCAGCGCAAGGTACTCTGAGT
CCCCICTGTAGTGCCTCTGCCAGACACACTGCCTGAGTTGAAGAGACACAGGCCACACATTTCAGGCTGGTTG
CCAGCGGACGTCAGCACTCACGGCCTGCGGGGACTCAGCACAGCTCTGGATTCTGGATCTCTCCTGCTGTAACCC
CACGCACAAGCCTGCAACCCCCAGAGCTCTTTGACAGGCTCCCAGGCCTCCCAGTCTGGACAAGCATGTGCAGT
CACGGGAGCTCAGCTCAGGCCAGGGCTGGGCTGTGCACCTGCCCTCCCACTGACCCAGACCCACTTCTCCAGAGA
GGCCTCTCTCCGCTGAGCTATTTCCCTTGCTAGTGTGCAGATATTTCCCTAACATGTCCTTTTTTGTATTTGTT
TGTACGGACCATAAATATAACTGTAGCTTTAAGACTAAAAA

919/6881
FIGURE 855

MSKRSWWAGSRKPPREMLKLSGSDSSQSMNGLEVAPPGLITNFSLATAEQCGQETPLENMLFASFYLLDFILALV
GNTLALWLFIRDHKSGTPANVFLMHLAVADLSCVLVLPTRLVYHFSGNHWPFGEIACRLTGFLFYLNMYASIYFL
TCISADRFLAIVHPVKSLKLRPLYAHLACAFLWVVVAVAMAPLLVSPQTVQTNHTVVCLQLYREKASHHALVSL
AVAFTFPFITTVTCYLLIIRSLRQGLRVEKRLKTKAVRMIAIVLAIFLVCFVPYHVNRSVYVLHYRSHGASCATQ
RILALANRITSCLTSLNGALDPIMYFFVAEKFRHALCNLLCGKRLKGPPPSFEGKTNESSLSAKSEL

920/6881
FIGURE 856A

GGGGAAGATGGCGGCTGCTCCTTTGGAGGAGCGGGATTGAGAGGATCGGGGTGGGGAGACCAACAAGAGAGACA
TTTCTGGCTCTGAAGGCGAACGCTTCGCTGGCCATTTAGGAGCTCTGCTCAAAGCCAGACGTATCCTAGAAGGAA
AACATCACCATGGCTACAGAAATTGGTTCTCCTCCTCGTTTTTTCATATGCCAAGGTTCCAGCACCAGGCACCT
CGACAGCTGTTTTATAAGCGACCTGATTTTGCACAACAGCAAGCAATGCAACAGCTTACTTTTGATGGAAAACGA
ATGAGAAAAGCTGTGAACCGAAAAACCATAGACTACAATCCATCTGTAATTAAGTATTTGGGAGAACAGAATATGG
CAAAGAGACCAGAGAGATATGCGGGCAATTCAGCCTGATGCAGGTTATTACAATGATCTGGTCCCACCTATAGGA
ATGTTGAATAATCCTATGAATGCAGTAACAACAAAATTTGTTCCGACATCAACAAATAAAGTAAAGTGTCCTGTA
TTTGTGTGTTAGGTGGACTCCAGAAGGAAGACGCTTGGTCACTGGAGCTTCTAGTGGGGAGTTTACCTGTGGAAT
GGACTCACTTTCAATTTTGAACAATATTACAGGCTCACGACAGCCAGTGAGGGCCATGACGTGGTCACATAAT
GACATGTGGATGTTGACAGCAGACCACGGAGGATATGTGAAATATTGGCAGTCGAACATGAACAACGTCAAGATG
TTCCAGGCACATAAGGAGGCGATTAGAGAGGCCAGTTTCTCACCACGGATAATAAATTTGCTACATGCTCTGAT
GACGGCACTGTTAGAATCTGGGACTTTCTTCGTTGCCATGAGGAAAGAAATCTCCGAGGGCATGGTGCTGATGTG
AAATGTGTAGACTGGCATCCAACCAAAGGGTTAGTTGTTTCAGGAAGTAAAGATAGTCAACAGCCAAATCAAGTTC
TGGGATCCCAAGACTGGGCAGAGTCTTGCAACACTTCATGCCCCATAAAAAACACAGTAATGGAAGTGAATTAAC
CTCAATGGCAATTGGCTACTCACAGCATCACGTGATCATCTCTGTAACTTTTTGATATCAGAAACCTAAAAGAA
GAGCTTCAAGTCTTCCGAGGTCATAAGAAAGAAGCCACAGCTGTGGCCTGGCATCCTGTTTCATGAAGGACTTTTT
GCCAGTGGAGGGTCTGATGGTTCTTTGTTATTCTGGCATGTTGGGGTAGAGAAGGAAGTGGGTGGGATGGAGATG
GCTCACGAAGGGATGATCTGGAGTCTGGCTTGGCATCCTCTTGGGCATATTCTCTGCTCAGGCTCAAATGACCAT
ACTAGCAAATTCGGACTCGAAACCGACCAGGTGATAAAATGCGAGATCGATATAATCTAAACCTTTTACCTGGA
ATGTCTGAAGATGGAGTAGAATATGATGACCTCGAACCTAATAGCCTGGCAGTAATTCCAGGAATGGGAATACCA
GAACAATAAAATTAGCTATGGAACAAGAACAGATGGGGAAAGATGAATCAAATGAAATTGAAATGACAATTCCA
GGTTTAGATTGGGAATGGAGGAAGTGATGCAAAAGGATCAGAAAAAAGTACCTCAGAAGAAAGTTCCTTATGCA
AAACCCATTCTCTCAGTTCAGCAGGCTTGGATGCAAAATAAAGTTCCAATTCTGCTCCAAATGAGGTGCTG
AATGACAGAAAAGAAGACATTAAATTGGAAGAGAAGAAAAAACACAAGCAGAAATTGAGCAAGAAATGGCTACA
TTACAATATACTAACCCACAACCTTCTGGAGCAACTTAAATTTGAAGACTTGCACAGAAACAAGTTGAGCAAATT
CAGCCTCCTCCCTCATCTGGCACCCCTCTCCTCGGACCCACGCTTTTCCAGGACAAGGTCCAATGTCTCAGATT
CCTCAAGGTTTTCAACAGCCCCATCCATCTCAGCAGATGCCAATGAACATGGCTCAAATGGGGCCTCCAGGTCCA
CAGGGACAGTTTAGGCCTCCTGGACCCAGGGACAAATGGGACCACAAGGTCTCCACTGCATCAGGAGGTTGGG
GGGCCACAAGGATTCATGGGACCACAGGGGCCCCAGGGCCCCGCCCAGGGGTTGCCACGGCCTCAGGACATGCAT
GGGCCCCAAGGAATGCAGAGGCATCCTGGACCTCATGGCCCTTTGGGACCTCAAGGGCCACCTGGACCACAAGGT
AGTTCTGGTCTCAAGGTCAATATGGGTCTCAGGGTCCACCTGGCCCCACAGGGTCACATAGGGCCCCAAGGCCG
CCTGGCCCTCAGGGTCACTTGGGCCACAGGGGCTCCGGGTACTCAAGGTATGCAAGGACCACCTGGTCCCAGA
GGAATGCAAGGGCCTCCTCATCTCATGGGATCCAAGGCGGACCAGGGTCTCAAGGGATCCAAGGTCTGTGTCT
CAGGGACCTCTGATGGGATTGAATCCAAGAGGAATGCAGGGGCTCCAGGCCCCGGGAGAACCAGGGTCTGCT
CCCCAAGGGATGATTATGGGCCACCCGCTCAAGAGATGAGAGGACCTCACCTCCAGGTGGACTACTGGGACAC
GGCCCTCAGGAAATGAGAGGTCTCAGGAGATCCGAGGCATGCAGGGGCTCCACCCCAAGGATCAATGCTGGGA
CCTCCCCAGGAATTGCGAGGGCCTCCAGGCTCACAAAGTCAGCAGGGGCCGCCCCAGGGCTCTTTAGGACCTCCA
CCCCAGGGTGGCATGCAAGGACCCCGGACCTCAGGGACAGCAGAACCAGCAAGAGGGCCACATCCATCTCAA
GGGCAATACCATTCCAGCAACAGAAAACGCCTCTGCTAGGTGATGGGCCCCGGGCCCCCTTCAACCAGGAAGGA
CAGAGCACAGGCCCCCACCCTGATACCAGGCCTAGGGCAGCAGGGAGCACAAGGTGCGATTCCCCCTCTGAAC
CCCGGACAAGGACCTGGCCCCAACAAAGTTTCAGAAGAGGAGCCCCGCGGAGGCATGAGGGCGTGCTCCCCCA
GAGGAAGGGATGGTTTTCTGGTCTGAAGACTTTGGTCCAGAGGAGAATTTTGATGCTTCTGAGGAAGCGGGCC
GAGGACGAGATCTCAGAGGTGAGGTGCGGGTACCCACAGGAGGAAGGAAGGGTTTACTTCCCACTCCTGACG
AGTTCCCTCGCTTTGAAGGAGGGCGGAAGCCAGATTCTGGGATGGAACAGAGAGCCTGGGCCAGGTCTGAGC
CTCTCAAGGCATGGACATGGCATCCCTACCTCCCCGAAAGCGCCCTGGCATGATGGGCCAGGCATCTGAGC
ACAGAGAGATGGAGGCCCCAGGAGGCCCTTCTGAAGACCGAGGAGGCAAGGCCGAGGGGGCCAGGACCTGCTC
AGAGAGTGCCCAAAATCTGGGCGTTCCAGTCTCTTAGACGGAGAGCACCACGATGGATACCACAGAGATGAACCTT

921/6881
FIGURE 856B

TTGGGGGCCCTCCAGGCAGTGGCACCCCTTCTCGAGGGGGCCGGAGTGGCAGTAACTGGGGTAGAGGGAGTAACA
TGAACCTCTGGCCCGCCGAGGCGAGGAGCTTCACGGGGTGGTGGAAAGGGTCCGGTAGAAGCTGGAACCTGAGTACCC
TGAGGCCTCTCTGGACAGTATGTAAGAACTTCTTGTGGACTCACCAAGAGAAACAAAAGGAAGCCTGCACCATTG
TAGCCCTGAACTCTTTTCTGGGCACCTGAATCCCAGGAACCCCTCAATGAGGTCTTCAAGATGAAGAGACTGCTGC
CAGCTACCAGCCTGGCTGGCCCTGTCTGTCCACCCTCATTGCCCAACTCCCATGTTGTTTTGTGAAGATAAAA
GCTGGGATCTTTTTCTTTTTTTAAGTCTCACAAGACATGGGGCATCTCCACAAATTAAGTTCCTGTCCATTG
GAAATTTGTTTCTATGTGTACAGTTTGTCTAGAGAAAAACAAAGTTTTTGTATGAATACAGAATGTGATTTACGCA
AGAATTGACAGAAAACCTAGTTGTGAAGTGCTTGCCTTAAGGAAACCTTTGGTTTTCCATCGCATCCCATCTGCCAA
GGAATGCACGGTTGTTGCTGATGTTTCATGAGCATATTAACCACTAGGTTATCTAAATTAATCTCAGCTGTGAAC
TTGTGTTTTTCAACACATTCTGTACACACATTTTTGTGACAGAAAAATACTTGAAAATTATTCTAATGTAGTTG
TAGTAGGTTCAAATTTTTATTAGACATGCTCTGACAGATGGAAAATTGTACTCTGAAGAGAAAATGCAATCTTC
TGGATGTCCAACGAGGGAGCTTCCCCCTTGAAGCATCTCCTAAGAAATCACAGTTTAGGTTTGAATCGCCTTGTTA
TGATGAGGAGAGGAACCTTGATGTTAGTCAGAAATAAAGTCACATGCCCTTTAGATAACTTGAAATCACACATGGA
GTTGATAAGGAATGGACCAAAATAGGTCCTCAAACAATTTTGCAACACTTAATTAAGTTTTCTCTCTTCTCTCT
CCATACTATGTGGAATAACAGTGTTACGGCAGGGCGTACTTCTAACTCACAGTAAGCCAGATAGATCTTGTTCA
TGGCAAATTCCTTGTTTAAATAAACATACCATTTAGCCAGTTTGGATTGGAAGCCAAAAATTCAGTTTGTTAT
ATACCCCATGATACTTTCTATACTGACCTTTATTTTGGTTTTACATGGAAGCTTTTATTAAAAGGACTGTCCTAT
GGCCACCTTAACATTTTATTACATTGGCAGCTTCAGTTTGGCATCAGTCAAATTTATTGGCAAATTTAGTGGT
GCTGGAAGTTTCTTTTTTCTTTTCTTTTTTGACAGAGTCTTGCTCTGTCAACCAGCCAGGAGTGCAGTGGTGC
AATCACGGCTCACTTCAGCTTTGACCCTGCCTCAGCTCCCTAGTAGCTAGGACTACAGGCATATGCCACCATGC
CCAGCTAATTTTTTGTGTGTGTGTTTTTGTAGAGATGGAGTTTACCACCTTTGTCTATGCTGGTTTCAAACCTC
TGGGCTCAAGCGGTCCACCTGCTGTGGCTCCCAAAGTGCTGGGATTGCAGATCTGAGCCACAGTACCTGGCCGC
TGGTGCTGGAAGTTTCAATGGTTGCTTAGAAAAGGCATACTGATATAGTTTAAACAGGATATGAAAATTTAAGTGG
TTCTCTATGAATGTAAAAAACAGCAAAAGCTAATTGATTTCTGTGGAGTGAGAAAATGTGAGCCATATACACTGC
TTTTCAGTGTTAGAATCTAGAGACTATTTTTTTTTTAACCTTGAAATTACAGCTTATTTAATCAGCGCTCCACCA
TAAACATGCTATTAGGAGGCTCAAGATTGCTTTAGGTTATGGGTATTTCTTCACTCACCCCACTGGTATAGCT
TAAAGAGAGAGAACTGTAATCAAATTTTCATGAAGCTAAAACCGGATCCAGAGACCAAAACCCATATCAAGATTAG
ACTTTTTTCCCTGAGATTCTAGTGGAAAAATTTTATTATTTTGGGCTTAACTTTAGAAACATAGTATCTCAG
TAAGCAGATAACCAGATCACATTTAAAAATTTTATTTTACCTTGAAGCTCTACTGGACTTGACAGATCACATTT
TGAGAGTGAATTCATTTAAACAATAAACCTCTACATTATACTTAT

922/6881
FIGURE 857

AGAACATCCAGTCACGGATAAAAAATGAGCTGGTTTCAGAAGGCCAAACTGGCCGAGCAGGCTGAGCAATATGATGA
CATGGCAGCCTGCATGAAGTCTGTAAGTAAGCAAGGAGCTGAATTATCCAATGAGGAGAGGAATCTTCTCTCAGT
TGCTTATAAAAAATGTTGTAGGAGCCCGTAAGTCATCTTGGAGGGTCTGCTCAAGTATTGAACAAAAACCGGAAGG
TGCTGAGAAAAACAGCAGATGGCTCGAGAACACAGAGAGAAAAATTGAGACGGAGCTAAGAGATATCTGTAATGA
TGTATTGTCTCTTTTGGAAAAGTCTTGATCCCCAATGCTTCACAAGCAGAGAGCAAAGTCTTCTATTGAAAAAT
GAAAGGAGATTACTACCGTTACTTGACTGAGGTTACTGCTGGTGATGACAAGATAGGGATTGTGGATCAGTCACA
ACAAGCATAACCAAGAAGCTTTTGAAATCAGCAAAAAAGAAATGCAACCAACACATCCTGTGAGATTGGGTCTGGC
CCTTAACTTCTGTGTTCTATTATGAGATTCTGAATCCCCAGAGAAAGCCTGCTGTCTTGCAAAGACCGCTTTTG
ATGAAGCCATTGCTGAACCTTGATACATTAAGTGAAGAGTCATACAAAGACAGCATGCTAATAATGCAATTACTGA
GAGACAACCTTGACATTGTGGACATCGGATACCCAAAGGAGACGAAGCTGAAGCAGGAGAAGGAGGGGAAAAATTAAC
CGGCCTTCCAACCTTTTGTCTGCCTCATTCTAAAATTTACACAGTAGACCATTGTGTCATCCATGCTGTCCCAAAA
TAGTTTTTTGTTTATGATTTATGACAGGTTTATGTTACTTCTATTTGGATTTCTATATTTCCCATGTGGTTTTTA
TGTTTAATATTAAGGGAGTAGAGCCAGTCAACATTTAGGGAGGTATCTGTTTTCATCTTGAGGTGGCCAATATTG
GGATGTGGAATTTTATACAAGTTATAAATGTTTGGCATAGTACTTTTGGTACATTGTGGCTTCACAAGGGCCAG
TGTAAGCTGCTCCATGTCTAAGCAAGAAAACTGCCTACATATTGGTTTGTCTGGTGAGAAATAAAGGGAT
CATTGATTCCAGTCAGAGGTGTAGTAATTGTGGGTACTTTTAAAGTTTGGAGCACTTTACAAGGCTGTGGTAGAA
ACATACCCCATGGATACCACATGTTAAACCATGTATATCTGTGGAATACTCAGTCTCATTGTGCACACCTTTGAC
TACAGCTGCAGAAAGTTTTCTTTAGATAAAGTTGTGACCCATTTTACTCTGGATAAGGGCAGAAACAGTTCACAT
TCCATTATTTGTAAAGTTACCTGCTGTTAGCTTTTCTATTATTTTGTCTACACTCATTTTATTGTATTTAAATGTT
TTAGGCAACCTAAGAACAAATGTAAAGTAAAGATGCAGTAAAAATGAATTGCTTGGTATTTCATTGCTTCATGTA
TATCAAGCACAGCAGTAAACAAAAACCCATGTATTTAACTTTTTTTTAGGTTTTTGTCTTTGTGATTTTTTTT
TGATACTTGCCTAACATGCATGTGCTGTAAAAATAGTTAAACAAGGAAATAACTTGAGATGATGGCTAGCTTTGTT
TAATGTCTTATGAAATGTTTCGTGGATAATCCAAGCATAATTGTTAAGAACACGTGTATTAAATTCATGTAAGTG
AATAAAAGTTTTATGAATGGACTTTTCAACTACTTTCTCTACAGCTTTTCTGTAAATTAGTCTTTTGGTTCTGA
AATTTCTCTAAAGGAAATTGTACATTTTGAATTTTATTCCTTATTCCTCTTGGCAGCTAATGGGCTTTTAGTA
AGTTTAAACAAAAATTTATCATAACAACAAAAATACTACTAATATAACTACTGTTTCCATGTCCCATGATCCC
GTCTCTTCCCTCCCAACCTGAAAAAATGAGTTTCTATTTTCTGGGAGAGGGGAGATTAAATGGAAAAAATG
TTAATATGTTCCATTTAAATTTTGGTATATGGCATTTCCTAACTTAAGAAGCCACAATGTTCTTGGCCCATCAT
GACAATGGGTAGCATTAACTATAAGTTTGTGCTTCCAAATCACTTTTCGGTTTTAAGAATTTCTTGATACTCC
TACAGCCTGCCTTCGATTTTGATCCTTTATCTGTTTGTGAGGTGCACAAGATTACCTTCCTTTTGTAGCCATCT
GTCTTGTCACCAACCATTCCTACTTGGTGGCCATGTACTTGGAAAAAGGCCGATGATCTTTCTGGCTCCACTCA
ATGTCTAAGGCACCTGCTTCCTTTGCTTGCTATCCACAGACTATTTCCCTCATCCTATTTACTGCAGCAATTGT
CTCCTTAGTTGACGAGATTGTGTTTATCTCCCTTTAAACCCCTACCTATCCTGAATGATCTGTCAATTGTCTGCCT
TTAAATCCTTCCTCTTCTCTCTATTCTCTAAATAATGATGGGGCTAAATTATACCCAAAGCTCACTTT
ACAAAAATTTCTCAGTACTTTGCAGAAAACACCAACAAAAATGCCATTTTAAAGAGGTGATTTTTTCTTT
TAGAATGTTAGAATGTAAGCTCCTCAAAAGCAGGGACAATGTTTCCGTATGTTCTACTGTGCTTAGTATACTGT
AAATGCTCAATAAATACTGATGATGGGAGGCAGTGAGTCTTGATGATAAGGGTGAGAAACCGAAATCCC

923/6881
FIGURE 858

GATGACATGGCAGCCTGCATGAAGTCTGTAAC TAAGCAAGGAGCTGAATTATCCAATGAGGAGAGGAATCTTCTC
TCAGTTGCTTATAAAAAATGTTGTAGGAGCCCGTAAGTCATCTTGGAGGGTCGTCTCAAGTATTGAACAAAAACG
GAAGGTGCTGAGAAAAACAGCAGATGGCTCGAGAACACAGAGAGAAAATTGAGACGGAGCTAAGAGATATCTGT
AATGATGTATTGTCTCTTTTGGAAAAGTTCTTGATCCCCAATGCTTCACAAGCAGAGAGCAAAGTCTTCTATTG
AAAATGAAAGGAGATTACTACCGTTACTTGACTGAGGTTACTGCTGGTGATGACAAGATAGGGATTGTGGATCAG
TCACAACAAGCATACCAAGAAGCTTTTGAAATCAGCAAAAAAGAAATGCAACCAACACATCCTGTCAGATTGGGT
CTGGCCCTTAAC TCTGTGTTCTATTATGAGATTCTGA ACTCCCCAGAGAAAGCCTGCTGTCTTGCAAAGACCGC
TTTTGATGAAGCCATTGCTGAACTTGATACATTAAGTGAAGAGTCATACAAAGACAGCATGCTAATAATGCAATT
ACTGAGAGACAAC TTGACAATGTGGTTTTTATGTTTAAATATTAAGGGAGTAGAGCCAGTCAACATTTAGGGAGGT
ATCTGTTTTCATCTTGAGGTGGCCAATATTGGGATGTGGAATTTTTATACAAGTTATAAATGTTTGGCATAGTAC
TTTTG

924/6881

FIGURE 859A

TTTTCGGCGCCGTGGCGGAATGGCGGCTTCCATCTCTGAGGCGAGCGACGCTATGGATCCCACAGTGGTTTGCTA
AGAAGGCCATTTTCAACTCTCCACTGGAGGCTGCTATGGCGTTCCCTCACCTGCAGCAGCCCAGCTTTCTACTGG
CTAGCCTGAAAGCTGACTCTATAAATAAGCCCTTTGCACAGCAGTGCCAAGACTTGGTTAAAGTCATTGAGGACT
TTCCAGCAAAGGAGCTGCACACCATCTTCCCATGGCTGGTAGAAAGCATTTTTTGGCAGCCTAGATGGTGTCCTCG
TTGGCTGGAACCTCCGCTGCTTACAGGGGCGCTGAATCCTGTGGAGTACAGCATCGTGATGGAATTTCTCGACC
CTGGTGGCCCAATGATGAAGTTGGTTTATAAGCTTCAAGCTGAAGACTATAAGTTCGACTTTTCTGTCTCTACT
TGCCTGGTCTGTGAAGGCGTCCATCCAGGAGTGCATCCTCCCTGACAGTCTCTGTACCACAACAAGGTCCAGT
TCACCCCTACTGGGGGCTTGGTCTGAACCTGGCCCTGAATCCGTTTCGAGTATTACATATTCTTCTTTGCCTTGA
GCCTCATCACTCAGAAGCCACTTCTGTGTCCCTCCACGTCCGTAATTCAGACTGTGCCTATTTTCCTCTGGTGG
ACAGGTACCTGTGATGGTTCTGCCCCACGAAGGCAGTGTGCCCCACCACTCTCTCCAGCCCAGGGGGGACCA
GCCCCCTACCACCTCCCAGGACACCAGCCATACCCCTTTGCTTCTATGGCCTCCACCACACTAGCCTCCTAAAGC
GACACATCTCTCATCAGACGTCTGTGAATGCAGACCCCGCTCCACGAGATCTGGAGGTGAGAACTCTGTCTCC
AGGTTTTTGTGAAATGTGGCTTCATCACTATTCTTGGAGATGTATCAAAAAATGCAGTCCCCTCATGCCAAGG
AGTCGTTACGCTACTGAGGAGCATGTGTTGGTGGTGCCTGCTGTGAAGCACCTGCACGCCTTTGCCAACA
GCCTGAAGCCAGAGCAGGCTTACCCTCCGCCCCACTCCACGCCACCAGCCCCCTGGAGGAGTTCAAACGGGCTG
CTGTCCCGAGGTTCTGTCCAGCAGAACTCTACCTCTTCTTGCAGCATTGCTTTGGCCACTGGCCCCCTGGACGCAT
CGTTCAGAGCTGTCTGGAGATGTGGCTGAGCTACCTGCAGCCGTGGCGGTACGCGCTGACAAGCAGGCTCCGG
GCAGCGACTCCAGCCCCGGTGTGTGTCGGAGAAATGGGCACCCCTTTGTCCAGGAGAACCTGTGATGTACACCA
AGTTGTTTTGTGGGCTTTCTGAACCGCGCTCCGCACAGACCTGGTCAGCCCCAAGCACGCGCTCATGGTGTTC
GAGTGGCCAAAGTCTTTGCCAGCCCAACCTGGCTGAGATGATTGAGAAAGGTGAGCAGCTATTCTGGAGCCAG
AGCTGGTCATCCCCACCGCCAGCACCGACTCTTACGGCCCCACATTCACTGGGAGCTTCTGTACCCCTGGC
CACCAGCGGTCACTGATGCCTCCTTCAAGGTGAAGAGCCACGTCTACAGCCTGGAGGGCCAGGACTGCAAGTACA
CCCCGATGTTTGGGCCCCGAGGCCCCGACCCCTGGTCTGCGCTCGCTCAGCTCATCACAGGCCAAACACACAG
CCAAGTCCATCTCCGACCAGTGTGCGGAGAGCCCGCTGGCCACTCCTTCTCTCATGGCTGGGCTTTAGCTCCA
TGGACACCAATGGCTCCTACACAGCCAACGACCTGGACGAGATGGGGCAAGACAGTGTCCGGAAGACAGATGAAT
ACCTGGAGAAGGCCCTGGAGTACCTGCGCCAGATATTCCGGCTCAGCGAAGCGCAGCTCAGGCAGTTCACACTCG
CCTTGGGCACCAACCAGGATGAGAATGGAAAAAGCACTCCCCGACTGCATCGTGGGTGAGGACGGACTCATCC
TTACGCCCCCTGGGGCGGTACCAGATCATCAATGGGCTGCGAAGGTTTGAATGAGTACCAGGGGGACCCGGAGC
TGCAGCCCATCCGGAGCTATGAGATCGCCAGCTTGGTCCGCACACTCTTTAGGCTGTCTGTGCCATCAACCACA
GATTTGCAGGACAGATGGCGGCTCTGTGTTCCCGGATGACTTCTCAGGAGCTTCTGTGCTACCACTCACAG
AACCTGGGCTGGCCAGCAGGCACCTGTGAGCCCTGTGGGGCGGAGGCAGGTGGCCGGCCACACCCGCGGCCCA
GGCTCAGCCTGCGCTTCTTGGGAGTTACCGGACGCTGGTCTCGCTGCTGCTGGCCTTCTTCTGTGGCTCTCTGT
TCTGCGTGGGGCCCCCTCCCATGCAGCTGTCTGCTACCCCTGGGCTATGTCTCTACGCTCTGCCATGACACTGC
TGACCGAGCGGGGAAGCTGCACACAGCCCTGAAGGTGTCAGCTGCCTTACAGCAGGCTGGAGGGATTGGCCACA
CAGCCCCACCCTTGGGCTGAGAGGACCTGGGAAGCCCTCCAGGAGGGAACACGGTCATCTCGGGCTTCTGGAG
CGGGGTTCTGTCAGCCGCAGAGGCATCTGGAGGAAACGCAACCAAGAAAGGAAGGCAGGTGGGCCCCAGCAAAGG
AGTAGCTGCCAGGGCTCAACAGCTACGCTCTGTGACAGCGCAGAGCTCAGCGGCGGCCCTTTCCCTCCCTCCGCCA
AGGACTCACGGCCAAGCCAGCTCTCGGGGCTTTTTTCCACTGCCCATTGGCTACTCTGCTGCACCAAGCTTGG
GAGCCAGCCTGCCAACAGCCACCTGGGCTGGCCTCCCCACTGGCTGGCCTTGAGGTTGGCAGAGTGGGTTGTGG
CGCTTCTCTCTGTGTGGGACAGGACAGTGGCTTAAGTCTCCACTCCAGGAAAGAAATCAAAGTTTCTAGAGT
TGTGAGAAAACAGAGAGTGGCTGTCTGATTCTTCACTGTGAGGGGCGTTCTTCACTGTTCTCCAGCTGTTCCA
AGACTGGGCGGTAGAATTCCATGTTTTCAGGAGCCTAAGACCCTCCAGAGCCAGGGGCTTACCGCAGACCCCA
AGCCATTGAGCACATACCCAAAGCAGTGGCCAACATCGCGGACCCCTGTGCCTTGTACAGATGGGTGCTGGTC
CTCAGGCGTTGGGGACACTGCTGGGTGATGGGTCGATTCTGCCAGTTTCTGCTGTGACGCAAGATGGTCA
GAAGCATTGTCACTTCACTAATCAAGTGTCAAAGACATGGCAACCGTTTCACTGGTACTTAAGTATTCAAAAT
ATACAACTACAGATTCTGTACAGAAACAGCAGGGGTCTTACCTTCACTTACCCACAGGCGACATGCGAGG
GAGAACAGCATCTCAGTGGTGATTTCAAACCAAGCCTTTGTTTTCGGTGTGGGGTTTTGGGGTTTTGCTTTAAT
GTTTTTGAAATGTAAATGTTGGGCTTTGTATTTGATGTAACTGAGCATAATGGCATTTTAGGGCTGTGACC

925/6881

FIGURE 859B

AAAATGAAGCTTGTAACGACCCATGGATCTGAATAAACATGTCCTTGCTTCTGAAAAAAAAAAAAAAAAAAAAA
AAAAAAA

926/6881

FIGURE 860

MAFPHLQQPSFLLASLKADSINKPFAQQCQDLVKVIEDFPAKELHTIFPWLVESIFGSLDGVLVGWNLRCLQGRV
NPVEYSIVMEFLDPGGPMMKLVYKLQAEDYKFDFPVSYLPGPVKASIQECILPDSPLYHNKVQFTPTGGGLNLNLA
LNPFEYYIFFFALSLITQKPLPVSLHVRTSDCAYFILVDRYLSWFLPTEGSPVPPPLSSSPGGTSPSPPPRTPAIP
FASYGLHHTSLLKRHISHQTSVNADPASHEIWRSETLLQVFVEMWLHHYSLEMYQKMQSPHAKESFTPTTEHVLV
VRLLLKHLHAFANSLKPEQASPSAHSHATSPLEEFKRAAVPRFVQQKLYLFLQHCFGHWP LDASFRAVLEMWLSY
LQPWRYAPDKQAPGSDSQPRCVSEKWAPFVQENLLMYTKLFVGFLNRALRTDLVSPKHALMVFRVAKVFAQPNLA
EMIQKGEQLFLEPELVIPHRQHLFTAPTFTGSFLSPWPPAVTDASFVKVSHVYSLEGQDCKYTPMFGPEARLTV
LRLAQLITQAKHTAKSISDQCAESPAGHSFSLSWLGFSSMDTNGSYTANDLDEMGOQSVRKTDEYLEKALEYLRQI
FRLSEAQLRQFTLALGTTQDENGKKQLPDCIVGEDGLILTPLGRYQIINGLRRFEIEYQGDPELOPIRSYEIASL
VRTLFRLLSSAINHRFAGQMAALCSRDDFLGSFCRYHLTEPGLASRHLLSPVGRROVAGHTRGPRLSLRFLGSYRT
LVSLLLAFFVASLFCVGPLPCTLLLTGLGYVLYASAMTLLTERGKLHQP

927/6881
FIGURE 861

GGCGCCGAGTAGCCGGGCCGGGCCGGAGCGCGGGCGGGCGGAGGCAGCTGCGCCCGCGCCTCCTGCCCTCCCA
GGCCCCGCGCCCCGCGCCCGGGCCCCGGCGATGGTGACACATGCGGCGGGCGGCGCGCCGGCGGCAGGACCATGGT
TGAGCGCGCCAGCAAGTTCGTGCTGGTGGTGGCGGGCTCGGTGTGCTTCATGCTCATCTTGTACCACTACGCGGG
CCCAGGACTGAGCCTGGGCGCGCCCCGGCGGCCGCGCGCCGCCGACGACCTGGACCTGTTCCCCACGCCCGACCC
CCACTACGAGAAGAAGTACTACTTCCCGGTCCGCGAGCTGGAGCGCTCGCTGCGCTTCGACATGAAGGGCGACGA
CGTGATCGTCTTCTGACATCCAGAAGACGGCGGGCACCACCTTCGGCCGCCACCTCGTGACAGACGTACGCCT
CGAGGTGCGGTGCGACTGCCGGCCCCGGCCAGAAGAAGTGCACCTGCTACCGGCCCAACCGCCGCGAGACTTGGCT
CTTCTCCCGTCTTCTCCACCGGTGGAGCTGCGGGCTGCACGCCGACTGGACCGAGCTACCAACTGCGTGCCCGG
CGTGCTGGACCGCCGCGACTCCGCCGCGCTGCGCACGCCCAGGAAGTTCTACTACATCACCTTGCTACGAGACCC
CGTGTCGCCGTACCTGAGCGAGTGGCGGCATGTGCAGAGGGGTGCCACGTGGAAGACGTGCTTGCATATGTGTGA
TGGGCGCACGCCACGCCTGAGGAGCTGCCGCCCTGCTACGAGGGCACGGACTGGTCGGGCTGCACGCTACAGGA
GTTTCATGGACTGCCCGTACAACCTGGCCAACAACCGCCAGGTGCGCATGCTGGCCGACCTGAGCCTGGTGGGCTG
CTACAACCTGTCTTCATCCCGAGGGCAAGCGGGCCAGCTGCTGCTCGAGAGCGCCAAGAAGAACCTGCGGGG
CATGGCCTTCTCGGCCTGACCGAGTTCACGCGCAAGACGCAGTACCTGTTTCGAGCGGACGTTCAACCTCAAGTT
CATCCGGCCCTTCATGCAGTACAATAGCACGCGGGCGGGCGGGCGTGGAGGTGGATGAAGACACCATCCGGCGCAT
CGAGGAGCTCAACGACCTGGACATGCAGCTGTACGACTACGCCAAGGACCTCTTCCAGCAGCGCTACCAGTACAA
GCGGCAGCTGGAGCGCAGGGAGCAGCGCCTGAGGAGCCGCGAGGAGCGTCTGCTGCACCGGGCCAAGGAGGCACT
GCCGCGGGAGGATGCCGACGAGCCGGGCGCGTGGCCACCGAGGACTACATGAGCCACATCATTGAGAAGTGGTA
GTGGCGGTGGTGGCCACGGGGAGGCCTCTTGGGGGTGTGGGGGATAAAACAGGACAGACGACAGGTCCACCCAA
GACTGTCAAGGGATGAGCATCCCAAACCTGCTCCACAGAGGTAGCTGCGTCCTGAAAAAAACAGAGCAGGGATG
TAGTGGGGCTGGGCAGGGATGGGGGCTTGAGAAATCAACAGGTGCAGCCAGTGGGTGAGGAAAGCGTGCTCG
AAGGATGCCATGGTCAGGGCAGGGCCTCCAGAGCAGGTGTTGTGCTGAGCTGCTCTCTGCTGCTCTTGGATT
TATCGCAAAACTGAAGGTTTGCGCAAGAGACGAGGACAGCGGAAAGTGGACCTGCCAGGCCGGGAGTGTGTCCC
TCACCAACTATGCACACAGCACTCGCTCTTAGCTCCTCTGTCCGGGCTACTAGGAGTGAGACCAGCTTCTGGCAA
CTGCCCCAGCTCCAGGCCATCCCATAGCCCCCTCCTTCTGGCTGCCCCCAATGCCCCGAGGCCTGGGGAGCCCC
CAGCTCACCCATCTGTAGCTCCCTCAAAGTCAGGGCCCAACCCATCTGAGGCAGAGAAGACTCGAGTCCAGCCCC
CAGGAAGCCTGCTCCCCTCTCTGGCCATGCTCCTGCTTCATGCTTTGGGTGAGGAGCCAAAGCTGATGTTTCAG
GCCCCACCACTCCCTACAGTCCTCAGACC

928/6881
FIGURE 862

GACTGCCCCGTACAACCTGGCCAACAACCGCCAGGTGCGCATGCTGGCCGACCTGAGCCTGGTGGGCTGCTACAAC
CTGTCTTCATCCCCGAGGGCAAGCGGGGCCAGCTGCTGCTCGAGAGCGCCAAGAAGAACCTGCGGGGCATGGCC
TTCTTCGGCCTGACCGAGTTCAGCGCAAGACGCAGTACCTGTTTCGAGCGGACGTTCAACCTCAAGTTCATCCGG
CCCTTCATGCAGTACAATAGCACGCGGGCGGGCGGCGTGGAGGTGGATGAAGACACCATCCGGCGCATCGAGGAG
CTCAACGACCTGGACATGCAGCTGTACGACTACGCCAAGGACCTCTTCAGCAGCGCTACCAGTACAAGCGGCAG
CTGGAGCGCAGGGAGCAGCGCCTGAGGAGCCGCGAGGAGCGTCTGCTGCACCGGGCCAAGGAGGCACTGCCCGCG
GAGGATGCCGACGAGCCGGGGCGCGTGCCACCGAGGACTACATGAGCCACATCATTGAGAAGTGGTAGTGGCGG
TGGTGGCCACGGGGAGGCCTCTTGGGGGGTGTGGGGGATAAAACAGGACAGACGACAGGTCCACCCAAGACTGTC
AAGGGATGAGCATCCCCAAACCTGCTCCACAGAGGTAGCTGCGTCTGAAAAAAACAGAGCAGGGATGTAGTGGG
GCTGGGCAGGGATGGGGGCTTGAGAAATCAACAGGTGCAGCCAGTGGGTTCAGAGGAAAGCGTGCTCGAAGGATG
CCATGGTCAGGGCAGGGCCTCCAGAGCAGGTGTTGTGCTGGAGCTGCTCTCCTGGCCTCCTTGGATTATCGCA
AAAAGTGAAGTTTTCGCAAGAGACGAGGACAGCGGAAAGTGGACCTGCCAGGCCGGGAGTGTGTCCCTCACCAA
CTATGCACACAGCACTCGCTCTTAGCTCCTCTGTCCGGGCTACTAGGAGTGAGACCAGCTTCTGGCAACTGCCCC
AGCTCCAGGCCATCCCATAGCCCCCTCTCTTCTGCTGCCCCCAATGCCCGGAGGCCTGGGGAGCCCCCAGCTCA
CCCATCTGTAGTCTCCCTCAAAGTCAGGGGCCACCCCATCTGAGGCAGAGAAGACTCGAGTCCAGCCCCCAGGAAG
CCTGCTCCCCCTCTCTGGCCCCATGGTCTGCTTCATGCTTTGGGTTCAGGAGGCCAAAGCTGATGTTTCAGGCCCCAC
CCACTCCCTACAGTCTCTCAGACCAAGGAGGGGTTTGGGTAGTAGGCCCGAGCTGCATTGCCGGCCTTCCTCGGGC
CAACTGGCAGCCAGGAGTGGGGAGGCTTTGGCCAGGGATGCTGCCACTTGTGCGTGAGTCCGCGGCTGGCCCTT
GGAGGTGACCATCCAGGCAGGCCTGGCTCAGACTGGAAGGGCTGGGGACCGAGGGCTCCCTGCCTCTGTTCTCC
TTTCTGACCCACTGGGATTTGTAGCAGGCTGCCCCAGCCCCATCACCGAAACACATACTCAAGAGCTCAAATAC
CACTGCTCCCAACAGCGTACGGATTAAGTTTATCAGGCTTCCATCGGCTGGAGCATGGGACCATAGCCCCCTGCCC
AGGAGCCGTACCCTCGGACCACAGTGGCTCTGTATGGCCAGGAACCTGGGACTCGAGCTTTCAGATTCTCAACTAG
CCTTGGCAAAACAGCTGTAGGTGGCCTCCCTGACAACAGACACTCAGCCCTCCCAACCTGGCTCTCCTTGCATT
TCCCCATGCTCCCCACCCCCCTGGCAAAAGGCTGGCCATGCTCTGTTCCCAAGCAGCCGCGCAGGTTTCCCCACTGG
CTGCAATGGCCCTACCAAAAGCCATGTTGCATATCCGTTGTAAGCACGTGCCCTGTGCCCTGTCCCCATTCTTA
TGCCCTAGGAGGCCAAGCTGGTGTCTTAGGAGGGCCACACAGGCACCCTGGATCCCCCAGAGAGTAGATTGGT
GTGCTCAGGCCGAGGCTGACTCAGAGGTAGGGCAGTGGGCTCTGCAGGCCACCTGGCTGGGGTGGTGGGGGTC
CTCTCTCCTGCCCCAGCTTCCACTCAGCCACCACAGTTGCCACACCATGGGGTGGAGACGTGGGTACCCACGGGC
TTGGGAGCAAGCGCCTTCTGCAGCACAGGAAGCCGAAGCTGGGGTCAGGTGAGGTGCTACCCCTGGAGGTCTGG
CATAAGGGCCCCACCTCAGGCTCTCTACACTGGCCCCATTTTACTTTGGGGTCCAAGGACAGGATGGTCAACAG
GGCAGGGTGGACAGCGTGCCAGCGCCGCGCAGGGCCACCTCCCTGGGTGGATGCATCACACTAAGGAAGTGAGTG
CCAAGGGGATTTAGTGGTGTGGTTCTTTCAAAGGGAGGTGAGGGTCAATGGGAATCTGCTCGGACACTCAACATG
GGGGTGGGTGCACTCCTTGGAGGAGGAGGAACACGTTGAGGGGATTGTGAGGTCTTGACAAGCCACGTGGGGCA
CCTTGGCTTCCCGGCAGGAGGTGGACACCCAGCCAGAGGCCTGGCTCAAGGTGACCTCACCTTACCATGGGCTT
CCTGGGTGCGCGGGCCTGAGCGCAGGTGTTTTGTACATATTGGAATATGTGTTAACTTATGCCCCGCATCCCAA
CTCACACGGAAGCACGGGTCTTGTCTCAGTCTCTTCGCTGCATTGGAAGCAGTCTCCTCTCGGGCCAGCGCCG
GGCTGAGGTGTCCAGAGGCGGCGGCAGCTGGCAGTGCCCTCAGCCCCCAAGTGTCCAGCCTGGCACTTCCCATTC
AGGCCACCTGCTTTGGGTCAACAGTTCCTTTGCCAGCAGCATCTCCTAAATTGTAAGGACTCTGTCCACCGGGC
CCTCCAGGGCTGTGAGGACGGAACAGGCAGGGAGTGGAGCTAACAGCTTAGTCACCAGGACCCCCAGACCTGC
AAACGTCCCCTCCTGGAAGGGGAAGCCAGGAACAGCAGAACTGCCACAAAAACAAGGCTGTGAACCTTTTCGGGAA
CTGGAAGTGTAACTTGAACCCAGGATTGTTTAAAGCTTTATTTATTTATAGCTTCTTCTTAAAAAAAAGTGT
GAAGAAATTTTTGGTTATTTCATACAAAAAATGAGTTGATGATGGAAGCAAGTCATAATCATCTAATTGTTT
TTGCTAGGTGAGAAATGAATGTTAGCTGATGAAATAAACCTGACAAGG

929/6881
FIGURE 863

MLADLSLVGCYNLSFIPEGKRAQLLLESAKKNLRGMAFFGLTEFQRKTQYLFERTFNLKFIRPFMQYNSTRAGGV
EVD E D T I R R I E E L N D L D M Q L Y D Y A K D L F Q Q R Y Q Y K R Q L E R R E Q R L R S R E E R L L H R A K E A L P R E D A D E P G R V P T E D
Y M S H I I E K W

930/6881
FIGURE 864

GGAAAAAGCGACTTGTGGCGGTCGAGCGTGGCGCAGGCGAATCCTCGGCGCTAAGCAAATATGGACCTCGCGGC
GGCAGCGGAGCCGGGCGCCGGCAGCCAGCACCTGGAGGTCCGCGACGAGGTGGCCGAGAAGTGCCAGAACTGTT
CCTGGACTTCTTGGAGGAGTTTCAGAGCAGCGATGGAGAAATTAATACTTGCAATTAGCAGAGGAACTGATTCTG
TCCTGAGAGAAACACATTGGTTGTGAGTTTGTGGACCTGGAACAATTTAACCAGCAACTTTCCACCACCATTCA
AGAGGAGTTCTATAGAGTTTACCCTTACCTGTGTGCGGCCTTGAAAACATTCTGTCAAAGACCGTAAAGAGATCCC
TCTTGCCAAGGATTTTTATGTTGCATTCCAAGACCTGCCTACCAGACACAAGATTCGAGAGCTCACCTCATCCAG
AATTGGTTTGCTCACTCGCATCAGTGGGCAGGTGGTGGGACTCACCCAGTTCACCCAGAGCTGTGAGCGGAAC
TTTTCTGTGCTTGGACTGTCAGACAGTGATCAGGGATGTAGAACAGCAGTTCAAATACACACAGCCAAACATCTG
CCGAAATCCAGTTTGTGCCAACAGGAGGAGATTCTTACTGGATACAAATAAATCAAGATTTGTTGATTTTCAAAA
GGTTCGTATTCAAGAGACCAAGCTGAGCTTCCTCGAGGGAGTATCCCCCGAGTTTAGAAGTAATTTTAAGGGC
TGAAGCTGTGGAATCAGCTCAAGCTGGTGACAAGTGTGACTTTACAGGGACACTGATTGTTGTGCTGACGTCTC
CAAGCTTAGCACACCAGGAGCACGTGCAGAACTAATTCCCGTGTGATGGTGTGATGGATATGAGACAGAAGG
CATTGAGGACTCCGGGCCCTTGGTGTAGGGACCTTTCTTATAGGCTGGTCTTCTTGCCTGCTGTGTTGCGCC
AACCAACCAAGTTTGGGGGAAAGAGCTCAGAGATGAGGAACAGACAGCTGAGAGCATTAGAACCAAAATGAC
TGTGAAAGAATGGGAGAAAGTGTGTTGAGATGAGTCAAGATAAAATCTATACCACAATCTTGTACCAGCCTGTT
CCCTACTATACATGGCAATGATGAAGTAAACGGGGTGTCTGCTGATGCTCTTTGGTGGCGTTCCTAAAGACAAC
AGGAGAAGGGACCTCTCTTCGAGGGGACATAAATGTTTGCATTGTTGGTGACCCAAGTACAGCTAAGAGCCAATT
TCTCAAGCAGTGGAGGAGTTCAGCCCCAGAGCTGTCTACACCAGTGGTAAAGCGTCCAGTGTGCTGGCTTAAC
AGCAGCTGTTGTGAGAGATGAAGAATCTCATGAGTTTGTGATTGAGGCTGGAGCTTTGATGTTGGCTGATAATGG
TGTGTGTTGTATTGATGAATTTGATAAGATGGACGTGCGGGATCAAGTTGCTATTGATGAAGCTATGGAACAGCA
GACCATATCCATCACTAAAGCAGGAGTGAAGGCTACTCTGAACGCCCCGGACGTCCATTTTGGCAGCAGCAAACCC
AATCAGTGGACACTATGACAGATCAAAATCATTGAAACAGAATATAAATTTGTGAGCTCCCATCATGTCCCCGATT
CGATCTCTCTTTATCCTTGTGGATGAATGTAATGAGGTTACAGATTATGCCATTGCCAGGCGCATAGTAGATTT
GCATTCAAGAATTGAGGAATCAATTGATCGTGTCTATTCCCTCGATGATATCAGAAGATATCTTCTCTTTGCAAG
ACAGTTTAAACCAAGATTTCCAAAGAGTCAGAGGACTTCATTGTGGAGCAATATAAACATCTCCGCCAGAGAGA
TGGTTCTGGAGTGACCAAGTCTTCATGGAGGATTACAGTGCAGACGCTTGAGAGCATGATTTCGTCTCTCTGAAGC
TATGGCTCGGATGCACTGCTGTGATGAGGTCCAACCTAAACATGTGAAGGAAGCTTCCGGTTACTGAATAAATC
AATCATCCGTGTGGAACACCTGATGTCAATCTAGATCAAGAGGAAGAGATCCAGATGGAGGTAGATGAGGGTGC
TGGTGGCATCAATGGTCAATGCTGACAGCCCTGCTCCTGTGAACGGGATCAATGGCTACAATGAAGACATAAATCA
AGAGTCTGCTCCCAAAGCCTCCTAAGGCTGGGCTTCTCTGAGTACTGCCGAATCTCTAACCTTATTGTGCTTCA
CCTCAGAAAGGTGGAAGAAGAAGAGGACGAGTCAGCATTAAAGAGGAGCGAGCTTGTTAACTGGTACTTGAAGGA
AATCGAATCAGAGATAGACTCTGAAGAAGAACTTATAAATAAAAAAAGAATCATAGAGAAAGTTATTTCATCGACT
CACACACTATGATCATGTTCTAATTGAGCTCACCCAGGCTGGATTGAAAGGCTCCACAGAGGGAAGTGAGAGCTA
TGAAGAAGATCCCTACTTGGTAGTTAACCTAACTACTTGCTCGAAGATTGAGATAGTGAAAGTAACTGACCAGA
GCTGAGGAAGTGTGGCACAGCACCTCGTGGCCTGGAGCCTGGCTGGAGCTCTGCTAGGGACAGAAGTGTCTG
AAGTGATGCTTCCAGGATTTGTTTTTCAGAAACAAGAATTGAGTTGATGGTCTATGTGTCACATTATCACAGGT
TTCATACCAACACAGGCTTCAGCACTTCCTTTGGTGTGTTTCTGTCCCAGTGAAGTTGGAACCAATAATGTGT
AGTCTCTATAACCAATACCTTTGTTTTCATGTGTAAGAAAAGGCCCATTACTTTTAAGGTATGTGCTGTCTATT
GAGCAAATAACTTTTTTTCAATTGCCAGCTACTGCTTTTATTCATCAAAATAAAATAACTTGTCTG

931/6881
FIGURE 865

MDLAAAEPGAGSQHLEVRDEVAEKCQKLFDFLEEFQSSDGEIKYLQLAEELIRPERNTLVVSFVDLEQFNQQL
STTIQEEFYRVYPYLCRALKTFVKDRKEIPLAKDFYVAFQDLPTRHKIRELTSSRIGLLTRISGQVVRTHPVHPE
LVSGTFLCLDCQTVIRDVEQQFKYTQPNICRNPVCANRRRFLDITNKS RFVDFQKVRIQETQAE LPRGSIPRSLE
VILRAEAVESAQAGDKCFTGT LIVVPDVSKLSTPGARAETNSRVSGVDGYETEGIRGLRALGVRDLSYRLVFLA
CCVAPTNPFRGGKELRDEEQTAESIKNQMTVKWEKVFEMSQDKNLYHNLCTSLFPTIHGNDEVKRGVLLMLFGG
VPKTTGEGTSLRGDINVCIVGDPSTAKSQFLKHVEEFSPRAYTSGKASSAAGLTA AVVRDEESHEFVIEAGALM
LADNGVCCIDEFDKMDVRDQVAIHEAMEQQTISITKAGVKATLNARTSILAAANPISGHYDRSKSLKQININLSAP
IMSRFDLFFILVDECNEVTDYAIARRIVDLHSRIEESIDRVYSLDDIRRYLLFARQFKPKISKESEDFIVEQYKH
LRQRDGGSGVTKSSWRITVRQLES MIRLSEAMARMHCCDEVQPKHVKEAFRLNKSII RVETPDVNLDQEEEIOME
VDEGAGGINGHADSPAPVNGINGYNEDINQESAPKASRLRGFSEYCRISNLIVLHLRKVEEEEDESALKRSELVN
WYLKEIESEIDSEELINKKRIIEKVIHRLTHYDHVLIELTQAGLKGSTEGSESYEEDPYLVNPNYLLD

932/6881
FIGURE 866

CCGGGATCTCGAGATAGCCGCAGCTCTCGCGATCTTTCTGGAGCCGCACCTCCACGCGGAGTCCGAGCGCGTGTG
CTGAGACCCAGGGTCGGGAGGGCGGAGACTGGGAGGGAGGGAGAAGCCCTTTGGCCTGCCTTACGGAAGCCTG
CGAGGGAGGGTGGTGTCCACTGCCAGTTCGGTGTCCCGATGCCAGCGCCAGCGCCAGCCGCAAGAGTCAGGAG
AAGCCGCGGGAGATCATGGACGCGGCGGAAGATTATGCTAAAGAGAGATATGGAATATCTTCAATGATACAATCA
CAAGAAAAACCAGATCGAGTTTGGTTCGGGTTAGAGACTTGACAATACAAAAAGCTGATGAAGTTGTTGGGTA
CGTGCAAGAGTTCATACAAGCAGAGCTAAAGGGAAACAGTGTCTTCTAGTCCTACGTCAGCAGCAGTTTAAATGTC
CAGGCTCTTGTGGCGGTGGGAGACCATGCAAGCAAGCAGATGGTTAAATTTGCTGCCAACATCAACAAAGAGAGC
ATTGTGGATGTAGAAGGTGTTGTGAGAAAAGTGAATCAGAAAATTGGAAGCTGTACACAGCAAGACGTTGAGTTA
CATGTTCAGAAGATTTATGTGATCAGTTTGGCTGAACCCCGTCTGCCCCCTGCAGCTGGATGATGCTGTTTCGGCCT
GAGGCAGAAGGAGAAGAGGAAGAGAGCTACTGTAAACCAGGATACAAGATTAGACAACAGAGTCATTGATCTT
AGGACATCACTAGTCAGGCAGTCTTCCGTCTCCAGTCTGGCATCTGCCATCTCTTCCGAGAACTTTAATTAAC
AAAGGTTTTGTGGAAATCCAACTCCTAAAAATTATTTAGCTGCCAGTGAAGGAGGAGCCAATGTTTTACTGTG
TCATATTTTAAAAATAATGCATACCTGGCTCAGTCCCCACAGCTATATAAGCAAATGTGCATTTGTGCTGATTTT
GAGAAGGTTTTCTCTATTGGACAGTATTCAGAGCGGAAGACTCTAATACCCATAGACATCTAAGTGTGTTGTT
GGTTTGGACATTGAAATGGCTTTTAATTACCATACCACGAAGTTATGGAAGAAATGCTGACACCATGGTACAA
ATATTCAAAGGACTTCAAGAAAGGTTTCAGACTGAAATTCAAACAGTGAATAAACAGTTCCCATGTGAGCCATTC
AAATTTTTGGAGCCAACCTCTAAGACTAGAATATTGTGAAGCATTGGCTATGCTTAGGGAAGCTGGAGTCGAAATG
GGAGATGAAGACGATCTGAGCACACCAATGAAAAGCTGTTGGGTCAATTTGGTAAAGGAAAAGTATGATACAGAT
TTTTATATTCTTGATAAATATCCATTGGCTGTAAGACCTTTCTATACCATGCCTGACCCAAGAAATCCCAAACAG
TCCAACCTCTTACGATATGTTTCATGAGAGGAGAAGAAATATTGTCAGGAGCTCAAAGAATACATGATCCTCAACTG
CTAACAGAGAGAGCTTTACATCATGGAATTGATTTGGAGAAAATTAAGGCTTACATTGATTTCCTCCGCTTTGGA
GCCCCCTCCTCATGCTGGTGGAGGCATTGGATTGGAACGAGTTACTATGCTGTTTCTGGGATTGCATAATGTTTCGT
CAGACCTCCATGTTCCCTCGTGATCCCAAACGACTCACTCCTTAAATTACACTTTGCCACTTAACCTCCAGTGTG
GATGACAGAGCGAGACCCTGCCTCAAAAAAAAAAAAAAAAAAAGAAAGCCACACTTATTCTTTTCAGTAACCTGC
TAGTGACACAGGCTGTACTTTAGGTACTTAAATATGCACTAGAATAAAATTTGCAAGGCCCTAAAATATCACTGTT
ATTTTTGGAGTAATTCAGTATAGGTTTCGTTTTAAAGAGATTTTTATAACTTCAGACATGCATCAGTAGGAAATAA
CTTGAGAAATTCATATGGTTATGTTACAAATTCATATTCTGTTACTACAGTAAACGTTAAGAGTTTTAAACAGTT
AAGATTGTACAATTTTTCTTCTTTTCTATATTACAAGGGCCCCAGTGTTAATGTCTTAGATTTTCAGTATTTGAA
CTTATTTTTTTAAATTCGTGTCATTGAGATAAGAATAATTCAGGTAGCATCTGAAATTTAATGAATGTATAATTG
GCATATCATGGAAAATTAACCAGAAAGTATCAGTTCTTAAAGTTATGCCTAGAAATTATGTAAAGCTAAACTAC
TGGTTAGAAAGTATTTCAGTGTAAATATTGTATTAATTTGTTAAATTCTAAACTTGAATTTCAATAAAATTTTAAAG
CT

933/6881
FIGURE 867

ATGACAAAGAAAAATGCCTGCATGGGCCTTGTTTTCCAGATTCAATCAAGGCTCAAAGCTTTATCTTGTCTTA
ATAGCACCTTGCACCGATAGAGCCCCGGCAAGGCTTTGCGGGCCTTAATTATTTCCCAAGATCGATGAGACCCCT
CAGCCCCCTGGGCAGCAGCATCCACGGCAGTTTTCCAGATGGATGAGCTGCGTCGAAAGCTGCAGAGCAGACAG
GGATGCAGACAAGGTCCAGCAGCCCCAGCAGAGCCTGCGGCCTGCGGGTTCTCGGGCGGATGGCACTGCAAAAT
TCAGCCCTTAATTCAATCAATATGTGCCTTCGAGGAGTTTATAGTCATGGTTTGGTATGGTTGGTATGTGATGCA
CACAAAGCAGAGACAGCTGCATGTAGAAGGGCAGGGGAGCATGGAGGCCCCAGTACAGCCAGGAGGCCCTGGAG
CTTCAGGCAGCAGGACATGGGTCTGGATCTGTAGTCAGAGGGGAGGTCAGGAAAGAGCTTGTACACCAGCAAAAG
ATTGGATCCTGTGGTTCTCAAACCTTATTAGTCCTGGAGCTGTGTTTCAAGGAAATTCTGCAGAAGAACAGCAG
TTCCCTTGCAAGGATGACCCAGGGCCCCCTGGTTCCAGCAGCTCCAAGATGAGATGGCAGAAAGGGAGAGGGGG
AACCAAGGAGAGGGAAGGAGGAAATATATTAGTGCAGCTGCCCAACGTCCCTCAAGGACAAGCACAAGGGGT
GGTGTGGACAAGCATAATGCAATTGGCCCCAGGGACAGGCCATAGACCCCACTTCCAAACCCCACTTCTCCTCACA
GCCCCAGGCCTGAAGGGAACCCGCTGGATGCAAGCCGAGGTACCCAGAGATGGAGGTGGACAGACAGTGTGGG
ATGGCCCTGCACCTGAACTCCCAGGCACTACCTTGCATTGTCTAGTTTCTAATTTTACACACACTGGGGAGAGG
TCTGCTGTGGGCCAGGCTGGCCACCAGCGCTGGGTAAGTAGGGACTGGCCTAGCCACAGGCCTCAAGGATGCCGA
AGTCTGGTGGGGGAGACAAGCATGGAGAGAGACAGAGATGGCAGAGGCCAGGACAGGAAGCCTCTGGGCATCCTG
GCTGAGGACAGGGTGGCACACCCAGAGGAGTGAATCGTGGAAATGCCGAAAGCTAAAGAGAAGATTCCCACCA
GAGATCACAGAGCAGAATGCAAAAGTATAAAATTGCCCAACCAACCGTCACTCACGGACCCCTTTGTCCCCTTC
TCTTCCCCTCCACCCCATAGAAGTTTGGCTGACAACCACGGGTTTATACCACATTTCACATATGGGAAGAATCT
GTGGCCCAGAGTGGTTACATTAGAGCACCTCTCCCCTGAGAGCCACACAAGGGCCCATACCACCTTCCAGTTCA
TGGCTCCACCTTTGTCTGTCTCACAACCTTCCCAGACCTGGCCTATACCCACATGTTATTAATAATCTTCTA
GATATGCCTTTAGGATTTGTGCAAAAGACACTTGAGATAAAGTATCTGTCTAGAACCAACCTGGGATGTAGCCAAA
GCTAAGGAATATGTCTGTCTGTGGAGAGCAGATGATCAATAGGTGGTTCCCAGCTAATGGAGTACTGGAGCA
AGGCTCATAGGCCTTTTCAAGTCCATCAAAGGAAGATAAGTCCCCACTCATTGTCCCTGGAGGAACTGTGAGGTC
TTCTTATATTTGCATGTATGTGCCAAAGAGAGACATGTAAAAACCAGAGCCCGGAGACATAGAGAAGATCATTCA
AAAGCACATGCCAAGCTAAAGACAATACCAGGGACAGGACCTCATCTCTGACTCTTGGTCCAGTTCTCCCTCCA
ACTCAGCAACAGCCTTTGGATTCCACTTGGGAAATATGGAAGTGAACAGCCAGAAGAAACCTTCCCTAACACT
GAAACCAGTGGTGAATTTGGTAAACGCCCTGCAGAAGATATGGAAGAGGAACAAGCATTAAAGATCTAGAAAC
ACTGATGAGATGGTTGAATTACCCATTCTGCTTCAGAGCAAGAATGCTGGAGCAGTACTGGAAAAGGAGGCAAG
AATATTAAGGCTCTCCATACAGACTACAACGCCAGCATTTCAGTTCTAGACAGCAGTGGCCCCAGCATATATTG
CATATCAGTGTGATATCGAAACAATTGGAGAAATCTGAAGAAAATCATCACTACCTTGGGAAGAGGGCCCGCAG
TTGCCATCACCCACTGCAACCAGCCAGCTCCCGCTCGAATCTGATGCTGTGGAATACTTAAATTACCAACAATAT
AAAGGAAGGGACTTTGACTGCGAGTTGAGGCTGTTGATTATCAGAGTCTAGCAGGAGGAATTATTGGGGAATGT
TGTCTCATTTTCACTGATAGAGTTGTTCTTATTGGAGGAAAACCCAATAGGGTTGTAGAGTGCATAAAGATCATC
CTTGATCTTATTTCTGAGTCTCCCATCAAAGGACATGCACAGCCTTATGATCCCAATTTTACGATGAAACCTAT
GATTATGGTGGTTTTACAATGGTGGTTGCTGACGACCGTGGACGCCAGGGGAGACCTCATGACCTATAACAGA
AAAGGGAGACCTGGAACCGTTACGACGGCGTGGTTGATTTTCAAGTGTGATGAACTTGGGGCTCTGCAATAGAT
ACATGGAACGCATCAGAATGGCAGATGGCTTATGAACACAGGGTGGCTCCAGATATGATTATTCCTAIGCAGGG
TGTATGGCTCATATGGTGTCTTGGTGGACCTATTATTACTACACAAGGAGCTTTGATCAAAATTGATGAGCCT
TTAGAAGGATCCGAAGATCGGATCATTACCATTACAGGAACACAGGACCAGATACAGAATGCACAGTATTTGCCG
CAGAATAGTGTGAAGCAGTATTCTGGAAAGTTTTCTAA

934/6881
FIGURE 868

MTKKNACMGLVFQIQSRLKAFILFLIAPCTDRARQGFAGLNYFPKIDETPQPPGQQHPTAVFQMDELRRKLQSRQ
GCRQGPAPPAEPAACGFLGRMALQNSALNSINMCLRGVYSHGLVWLVCDAHKAETAACRRAGEHGGPSTARPLE
LQAAGHGSGSVVRGEVRKELVHQKIGSCGSQTLISGAVFQGNAAEQQFPCKDDPRAPGSQQLQDEMAERERG
NQGEGRRKYIQSCPNVPQGGQAQRGGVDKHNAIGPGTGHRPHFQTPFLLTAQGLKGTWRMQSRGHPMEVDROCG
MALHPETPRHYLALSVSNTHTGERSAVGQAGHQRWVSRDWP SHRPQGCRSLVGETSMERDRDGRGQDRKPLGIL
AEDRVAHPPEESESWKCRKLKRRFPPEITEQNMQKYKIAQPPSLTDPFVPFSSPPPHRSLADNHGFIPHFHIWEES
VAQSGYIRAPPPLRATQGP IPPSSSWLHLCPALTTFPDLAYTHMLLNLLDMPLGFAAKTLEIKYLSEPTWDVAK
AKEYVLFCEQMINRWFPANGVTGARLIGLSVPSKEDKSPLIVPGGNCEVFLYLHVCAKERHVKTRARRHREDHS
KAHAKLKTIPGTGPHLLTLGPVLPPTQQTAFGFHLGNMETEQPEETFPNTETSSEFGKRP AEDMEEQAFKRSRN
TDEMVELPILLQSKNAGAVTGKGGKNIKALHTDYNASISVLDSSGPQHILHISADIETIGEILKKIITLLEGPQ
LPSPTATSQPLESDAVEYLNYYQYKGRDFDCELRLLIHQSLAGGIIGECPPHFTDRVVLLIGGKPNRVVECIKI
LDLISESPIKGAQPYDPNFYDETYDYGGFTMVFAADDRGRPGDLMTYNRKGPRGNRYDGVVDFSADETWGS
TWNASEWQ MAYEPQGGSRDYDYSYAGCHGSYGD LGGP IITTTQ GALIKIDEPLEGSEDRIITITGTQDQIQNAQYLP
QNSVKQYSGKFF

935/6881
FIGURE 869

ATGAAGCAGTTAATTACCCCTGCTAAACAGCAGGAAGCATTGAGCAAAAACTCTGCTCACCTTCCTCAATCAGA
GGACATTTCTGCTCATCAGTCAAGCAGGCAGACCACAGGACTCTCAGACAAAGGCTGAGAGGTAGACTGATGAGG
AGGAGACTGTCAGTGAGCACAAAGGCCTTAAGATTAGCGACTGAAGGCTCCTGGAATGAGTATGTGGTTAGAGAC
TGGAACTACTCAGGAGTTTCCCTAGGTCTGCTCCAAATAGCACAAAGCAATGGATGATGAAGGATCCATGTTAATA
GCAGTGGATCCACAGTGCAGATTTTCAGGTGATCTGCTCCCTACGACTCACTAATTGGAGTCCCATTGGCCCCC
CATGTGACCCCTCTTGGATAAAAATCAAAATGGCTCTCGTCTTGTTTGTCTCCTGTTTGGCCCCAGTCAGGGCT
GGCACAGAGTGGCAGCTCTCTGCTCTCCACAGGGCTCCCCGCTCCACCCAGCCCGATAAAGCATGCCGACTGGGC
TACAAAGCCAAGCAAGGTTACATTATATATAGGATTTGTGTTCCGCGTGAGGAGCAAGCTGGACACCACTGTGGG
GCTCTAAGAGTCCTGAATTCGTACTGGGTTGGTGAAGATTCCACATACAAATTTTTTGAGGTTATCCTCATTGAT
CCATTCCATAAAGCTATCAGAAGAAATCCTGACACCCAGTGGATCACCAAACAGTCCACAAGCACAGGGAAATG
CGTGGGCTGACATCTGCAGGCTGA

936/6881

FIGURE 870

MKQLITPAKQQEAFEQKLCSPSSIRGHFCSSVKQADHRTLQRLRGRLMRRRLSVSTKALRLATEGSWNEYVVRD
WNYSGLVSLGLLQIAQAMDEGSMLIAVDPQCRFSGDLLPYDSLIGVPLAPHVTLLDKNQNGSRLVCFSCLPVRA
GTEWQLSALHRAFRSTQPDKACRLGYKAKQGYIIYRICVRREEQAGHHCGALRVLSYWVGEDSTYKFFEVLID
PFHKAIRRNPDQTQWITKPVHKHREMRGLTSAG

937/6881

FIGURE 871

ATGTGGGGCGACCTCTGGCTCCTCCCGCCTGCCTCTGCCAATCCGGGCACTGGGACAGAGGCTGAGTTTGAGAAA
GCTGCAGAGGAGGTTAGGCACCTTAAGACCAAGCCATCGGATGAGGAGATGCTGTTCATCTATGGCCACTACAAA
CAAGCAACTGTGGGCGACATAAATACAGAACGGCCCGGGATGTTGGACTTCACGGGCAAGGCCAAGTGGGATGCC
TGGAATGAGCTGAAAGGGACTTCCAAGGAAGATGCCATGAAAGCTTACATCAACAAAGTAGAAGAGCTAAAGAAA
AAATACGGGATATTGAGAGACTGGATTGTTTACTGTGCCATGTGTTTATCCTAAACTGAGACAATGCCTTGTTTT
TTTCTAATACCGTGGATGGTGGGAATTCGGGAAAATAACCAGTTAAACCAGCTACTCAAGGCTGCTCACCATACG
GCTCTAACAGATTAGGGGCTAAAACGATTACTGACTTTCCTTGAGTAGTTTTTATCTGAAATCAATTAAAAGTGT
ATTTGTTACTTT

938/6881
FIGURE 872

MWGDWLWLLPPASANPGTGTEAEFEKAAEEVRHLKTKPSDEEMLFIYGHYKQATVGDINTERPGMLDFTGKAKWDA
WNELKGTSKEDAMKAYINKVEELKKKYGI

939/6881
FIGURE 873

GTTTAGGGGCTGGTACAAAGGATGAATTGCACATTGTTGAAGTAGAGGCAATGAATTACAAAGGCAGTCCAATTA
AAGTAACACTGGCAACTTTGAAAATGTCTGTACAGCCAAAGGATTCCCTTGGAGGCTTTGAAATCACACCACCAG
TGGTCTTACGGTTGAAGTGTGGTTCAGGGCCAGTGCATATTAGTGGACAGCACTTAGTAGCTGTGGAGGAAGATG
CAGAGTCAGAAGATGAAGAGGAGGAGGATGTGAAACTCTTAAGTGTATCTGGAAAGCGGTCGGCCCGTGGAGGTG
GTAGCAAGATTCCACAGAAAAAAGTAAAGTTGCTGCTGATGAAGATGAAGATGATGATGATGAGGATTTTGATG
ATGGGGAAGCTGAAGAAAAAGCGCCGGTGAAGAAATCCATACGAGATACTCCAGCCAAAAATGCACAAAAGTCAA
ATCAGAATGGAAAAGACTCAAAACCGTCAACACCAAGATCAAAAGGACAAGACTCCTTCAAAAAAACAGGAAAAA
ACTTCTAAACACCGAAAGGACCTAGTTCCATAGAAGACATTAAAGCGAAAAATGCAAGCAAGTATAGAAAAAGCG
CATTGAACAGTCCTGGGCACTGCTGGTAAATTAAGCCCAAAGATGGGGAGAAAGGAAAAGGAGAGACAAATATAG
TCCATACTGAG

940/6881
FIGURE 874

MNYKGSPIKVTLATLKMSVQPKDSLGGFEITPPVVLRLKCGSGPVHISGQHLVAVEEDAEESEDEEEEDVKLLSVS
GKRSARGGGSKIPQKKVKVAADEDEDDDDDEFDDGEAEKAPVKKSIRDTPAKNAQKSNQNGKDSKPSTPRSKGQ
DSFKKTGKNF

941/6881
FIGURE 875

AGCGGGTTTGCGGGAGCGCCGCGTGGTTAGCGTCGGCGGCTTTTGGCATGCGGACTTTTTCTGGCCCGGCTGGGC
CAATCCTGTGCGTTAATCCGCAGGAAGATGTGAGTTTCAAAAGGAGGTGGCGCAGGTTTCGAAGCGCATAACCC
AGCGAAAAAACAAGAACAACTTACTCCTGGAGTAGTCTATGTGCGCCACCTACCTAACCTACTTGACGAAACCC
AGATCTTTTCATATTTCTCCAGTTTGGCACTGTGACACGGTTTCAGGCTGTCCAGAAGTAAAAGGACTGGAAATA
GCAAAGGCTATGCATTGTGGAGTTTGAGTCTGAGGATGTTGCCAAAATAGTTGCTGAAACAATGAACAACCTACC
TGTTTGGTGAAAGACTCTTGGAGTGTCATTTTATGCCACCTGAAAAAGTACATAAAGAAGCTCTTTAAAGACTGGA
ATATTCCATTTAAGCAGCCATCATATCAATCAGTGAAACGGTATAATCGGAATCGGACACTAACACAAAAGCTAC
GGATGGAGGAGCGAATTTAAAAAGAAAGAAAGATTACTCAGGAAGAAATTAGCTAAAAAGGAATTGACTATGATT
TTCCTTCTTTGATTTTACAGAAAACGGAAGTATTTCAAAAATAATCGTCAGACGTCTACAAAAGGCCAGGTTT
TACGTAAGAAGAAGAAAAAGTTTCAGGTAAGTCTTGACACTCCTGAGAAGACTGTGGATAGCCAGGGCCCCACAC
CAGTTTGTACACCAACATTTTTGGAGAGGCGAAAATCTCAAGTGGCTGAAGTGAATGATGATAAAGATGATG
AAATAGTTTTCAAACAGCCCATATCCTGTGTAAGAAGAAATACAAGAGACTCAAACACCTACACATTCACGGA
AAAAAGACGAAGAAGCAGCAATCAGTGATTTTCAATGTATTATTTCTTTGAAAAATATAATATTTTTATGA
GAGTGGACTTTGTATTTCACTAGGTACAATGGAATACAACCTTTGACAAGATTTTCAGAGGAAAAATACACTGTT
TGGTCAAGTTAAGGAAAGCAGTGTGTAATTTTGGATTGCCCTTGGCTGAAATACAGGGGTGCATACCATCT
TGCAGTGGCTTGGCTGACATTGCCTCTTTGTCTGGCCTCTAGTTTTCTTTTGATATTTTCATAGCTCTCCTTAGT
TTACTCTGCCTGGATAGAAAGTTGACCACTAACTGCAGGTTAAGTACTAACTGCAGCCTTTTCTGTGCCAGC
AATTAAAGACCACCAATCTTGTGTTGTCCATCTACATGGTTTGTGCGGGACATTTAACTCATGGAGGTGCTTTAGA
TTTCAACATCAGATGGTTGAAGCTGGAAGTTTAATTATATGTAGAGTGAGAAGGCAGTTCCAGTTTTAGCACAGA
TTTGTTTATGTGTTTCAAGTTTTAATAGAGATTCAAAAATGACTCATTTTTACCAATAATGTTAAATTAGTTTTGG
TTGTGCTAGCATGAATTAATAACCACCATTTTATACCAGTATCATCAGTGAAGAATTGTATTTCAAGATTCAAAC
AATAACCAGCAATTAACCTTTTTTCTACAATGTATTTGTTTTCGAGTAGGACTTGGGAGTCATTGGAAAAA
ATAATAAATTTTCCCCTTCATTAACAAAAA

942/6881

FIGURE 876

MATFSGPAGPILSLNPQEDVEFQKEVAQVRKRITQRKKQEQLTPGVVYVRHLPNLLDETQIFSYSQFGTVTRFR
LSRSKRTGNSKGYAFVEFESEDVAKIVAETMNNYLFGERLLECHFMPPEKVHKELFKDWNIPFKQPSYQSVKRYN
RNRTLQKLRMEERFKKKERLLRKKLAKKGIDYDFPSLILQKTESISKTNRQTSTKGQVLRKKKKKVSGLDTPE
KTVD SQGPTPVCTPTFLERRKSQVAELNDDDKDEIVFKQPI SCVKEEIQETQTPTHSRKKRRRSSNQ

943/6881
FIGURE 877

CCGCGGCGTCCACACTCGCCGCGCGCGCGCGCGCGCGGCTGGACCTTGCTGGCCCGCGGCGCCATGAGCCGCAGC
CTGGACTCGGCGCGGAGCTTCTGGAGCGGCTGGAAGCGCGGGCGGCGGGGAGGGGGCAGTCCTCGCCGGCGAG
TTCAGCGACATCCAGGCTGCTCGGCCGCTGGAAGGCTGACGGCGTGTGCTCCACCGTGGCCGGCAGTCGGCCA
GAGAACGTGAGGAAGAACCCTACAAAGACGTGCTGCCTTATGATCAGACGCGAGTAATCCTCTCCCTGCTCCAG
GAAGAGGGACACAGCGACTACATTAATGGCAACTTCATCCGGGGCGTGGATGGAAGCCTGGCCTACATTGCCACG
CAAGGACCTTGCCTCACACCCTGCTAGACTTCTGGAGACTGGTCTGGGAGTTTGGGGTCAAGGTGATCCTGATG
GCCTGTCGAGAGATAGAGAATGGGCGGAAAAGGTGTGAGCGGTACTGGGCCAGGAGCAGGAGCCACTGCAGACT
GGGCTTTTCTGCATCACTCTGATAAAGGAGAAGTGGCTGAATGAGGACATCATGCTCAGGACCTCAAGGTCACA
TTCCAGAAGGAGTCCCGTTCTGTGTACCAGCTACAGTATATGTCCTGGCCAGACCGTGGGGTCCCCAGCAGTCCT
GACCACATGCTCGCCATGGTGGAGGAAGCCCGTCCGCTCCAGGGATCTGGCCCTGAACCCCTCTGTGTCCACTGC
AGTGCGGGTGTGGGCGAACAGGCGTCTGTGCACCGTGGATTATGTGAGGCAGCTGCTCCTGACCCAGATGATC
CCACCTGACTTCAGTCTCTTTGATGTGGTCTTAAAGATGAGGAAGCAGCGGCTGCGGCCGTGCAGACAGAGGAG
CAGTACAGGTTCTGTACCACAGGTGGCTCAGATGTTCTGCTCCACACTCCAGAATGCCAGCCCCACTACCAG
AACATCAAAGAGAATTGTGCCCCACTCTACGACGATGCCCTCTCTCTCCGACTCCCCAGGCACTTCTCGCCATA
CCCCGCCCACCAGGAGGGGTCTCAGGAGCATCTCTGTGCCCGGGTCCCCGGGCCACGCCATGGCTGACACCTAC
GCGGTGGTGCAGAAGCGCGGGGCTCCAGCGGGCGCGGGAGTGGGACGACAGACGGGGACGGGGACGGGGACGGGG
GCGCGCAGCGCGGAGGAGGCGCGCTCTACAGCAAGGTGACGCCGCGCGCCAGCGACCCGGGGCGCACGCGGAG
GACGCGAGGGGACGCTGCCGTGGCCGCGTTCCTGTGACCAAAGTCTGCCGATCTGGCGCTACGAGGACGTG
GCGGGTGGAGCTCAGACCGGTGGGCTAGGTTTCAACCTGCGCATTTGGGAGGCCGAAGGGTCCCCGGGACCCGCT
GCTGAGTGGACCCGGGTGTAAGTCTAACGCCAGTTCCTGCCTGTTGCTCTTGTGAGCTCGGACTGCTGATGCCC
CGGTGCTGCTGAGCGCCGTGCGCAGAATGGAACAGTGGGCTGGATCAAAGTTAAAGTTTCTCAGGGTGGGAAA
TGTGGGGGCTTTGCCCAATGACTGTAGCATTCAAGGCTTGAGGCTGGAGGAGGTAGCTAGGGTATAGTGGCTGG
TGAGGCTGCACAGAGCAGATTCAAGAAAGAAGATCAGGAAGGGCATGACCCCTGAGTTATGAAGGGGAGAAGGG
ACAGATGAGCTTCCGGAGACTGCTCTCTCTCACCACACAGCACTAGTCCATCCTCAGCACCTGAGCCTCCCTCACT
TGGACACTCAGGGGACCACACAGAGAAGTGGATGGACACTTCGCCATCCAGGCAGAACTAAGCCAGGCATAACCA
CAGCCAAGCAGATTAACCCAGGCAGACCGATAAAAAGACCTCCAGATAGGCAGACAGACAGATGGACCACCAAC
CTGGACAGACAGCCAAAGCTTCAGAGATACAGTCCACAGGTGGACAAAGGGATCCCCAGCCAGAGAGAGAGAC
CAGCCAACAGCTTGATAGACAGTGCAGCCAGAGAGACCACCAACAGAGCCCCCAAAGACAGACATCTCTGCT
AGCTGGACAGCCAGGTGGACCCCTAAGTTAGATTACTAGACAGATATAAACAGATCCCTGCTGAACAGATACA
CAGAGTTCTCAGACCCACCCCAACCTCAGGTGGGCTGGCTGGCTGACAGACCTTCTGGCCAGACAGACTCCTA
ACCAACCAGATGGACTGCCAGACAGGCAGACATCAGCCACATGGAATCCTGACATCCAGCCAGCCGGCCAGACT
CTCATCTTGATGTCTTGATGGATGGACCCAGCTAGTCAGACATGATCCTCCAGATTGACAGACAAGTCCCCCAA
ATGAGTACACATCTCCAGCTATTACAGACAGATGGACCCCAAGTATCAGGACCTATCTAGGCAGACCCAGCCA
GACCCCGCCAGACAGACTCCCAACCAGACTGACCCCTTACTATTACACAGCCTGCCGAGTAGCTGGGACTACA
GGTCTAATTTTTTTTTTTTTTAAAGAAATGAGTTTTTGCCATGTTGCCAGACTGGTCTTGAATCCCAACCTCAA
GCAATCCTCCTGCCTCAGCCTCCCAAGTGCTGAGATTACAGGTGTGAGCCACCAGGCTCAGCCCCCTAAGATT
GAAACACTTTAAATGGCCCATGGTAGGGTCTCTGCTAGGATAAAACATTAAAGCGGCTGTTAAAGAAATAAAGG
AGGACACGTCTCTGTGC

944/6881
FIGURE 878

TGGAAGCGGCACTCAAGATGCTGCGCCGCGAGGCCCGCCTGCGCCGCGAGTACCTGTACCGCAAGGCCCGGGAGG
AGGCGCAGCGCTCAGCCCAGGAGAGGAAGGAGCGGCTGCGGCGCGCGCTGGAAGAAAACCGCCTGATTCCCCTG
AGTTACGCCGAGAGGCTCTGGCCTTACAGGGGTCCCTGGAGTTTGATGATGCTGGAGGTGAAGGTGTGACCAGCC
ACGTGGATGATGAATACCGATGGGCAGGAGTCGAGGATCCCAAGGTTATGATCACTACCTCCCGAGACCCAGTT
CCCGCCTCAAGATGTTTGCAAAGGAGCTGAAGCTGGTGTTCGCCGGCGCCCAGCGAATGAACCGAGGTGACATG
AAGTGGGGGCACTGGTGGAGCCTGCAAAGCCAACGGCGTCACCGATCTGCTGGTCGTTACGAGCATCGGGGCA
CACCTGTGGGGCTCATCGTCAGCCACCTGCCCTTTGGTCTACTGCCTACTTCACGCTGTGCAATGTGGTCATGC
GGCATGACATCCCAGACCTGGGCACCATGTGCGAGGCCAAGCCCCACCTCATCACACAGGCTTCTCCTCCCGCC
TGGGCAAGCGGGTCTCTGACATCCTCCGATACCTATTTCCCGTGCCCAAAGATGACAGCCACCGGGTCATCACCT
TCGCAAACCAGGACGACTACATATCATTCCGGCACCATGTGTATAAGAAGACAGACCACCGCAACGTGGAGCTCA
CTGAGGTGCGGCCCCGCTTTGAGCTGAAGCTGTACATGATCCGTCTGGGCACGCTGGAGCAGGAGGCCACAGCAG
ACGTGGAGTGGCGCTGGCACCCCTTACACCAATACCGCACGCAAGAGAGTCTTCCTGAGCACCGAGTCAGCACACT
CACCCTCAGTCAGGACATGGACTTGGAATCAGGATGGGGCTGTCATAGACAGACCCACCAGTAGGAACTGTCA
CAGAAATGGCCTGCTGAATGCGGATGTGGAATGTGGCGGGTGGAGAGGTCTGAATAAACCGTCTGTGTCATGGCA
AAAAAAAAAAAAAAAAAAAAA

945/6881
FIGURE 879

MLRREARLRREYLYRKAREEAQRSAQERKERLRRALEENRLIPTELRREALALQGSLEFDDAGGEGVTSHVDDEY
RWAGVEDPKVMITTSRDPSSRLKMFALVFPVGAQRMNRGRHEVGALVRACKANGVTDLLVVHEHRGTPVGLI
VSHLPFGPTAYFTLCNVVMRHDIPDLGTMSEAKPHLITHGFSSRLGKRVSILRYLFPVPKDDSHRVITFANQDD
YISFRHHVYKKTDRNVELTEVGPRFELKLYMIRLGTLEQEATADVEWRWHPYTNARKRVFLSTE

946/6881
FIGURE 880

GGCGGCGGCAGCGGGTTCGGTTGCGCGTGACGCACGGGGTGGGAGCGGAGCCCAGGCCGGGAGCAGGCGCCTCTG
CCAGAGACCATGGGGAACGTGTTGGCCGCCAGGTGCGCGCCCGCAGGGCCGCCACCGCCGCCTGCGCCGGCCCTC
GTGGAGCTGCCGCCACCTCCGCCCTCGCCGCCGGGCTTCACGCTGCCGCCGCTGGGAGGCGGCCTGGGCGCCGGC
ACCAGTACGAGTCGAGGTTCCGGAAGTACCCCCGGGGCTGCAACTGCCAGCGCCTCAGGGGCGCCGAGGATGGG
GCCTGCGGCTGCCTGCCCAACCCGGGCACATTCGGGGAGTGCCACCGGAAGTGCAAGGAGCTGTTTCCCATTTCAG
ATGGAGGGTGTCAAGCTCATAGTCAACAAAGGGTTGAGTAACCATTTCCAGGTGAACACACAGTAGCCCTCAGC
ACAATCAGGGAGTCCAGCTACCACTTCGGGGTCACGTATGTGGGGACAAAGCAGCTGAGTCCCACAGAGGCGTTC
CTTGATTGGTGGGTGACATGGACAACAGCGGCAGTCTCAACGCTCAGGTCATTACCCAGCTGGGCCCCCGCCTC
AGGTCCAAGATGGCCATCCAGACCCAGCAGTCGAAGTTTGTGAACTGGCAGGTGGACGGGGAGTATCGGGGCTCT
GACTCACAGCAGCCGTAACCCCTGGGGAAACCCAGACGTCCTCGTGGGTTTCAGGAATCCTCGTAGCCCACTACCTC
CAGAGCATCACGCCTTGCCTGGCCCTGGGCAGAAATGCTGGTCTACCACCAGCAGCCTGGGGAGGAGGGCACTGTC
ATGTCTCTAGCTGGGAAATACACATTGAACAACCTGGTGGCAACGGTAACGTTGGGCCAGGCGGGCATGCACGCA
ACATACTACCACAAAGCCAGTGACCAGCTGCAGGTGGGTGTGGAGTTTGAGGCCAGCACAAGGATGCAGGACACC
AGCGTCTCCTTCGGGTACCAGCTGGACCTGCCCAAGGCCAAAGGCTCTGTGGATAGCAACTGGATCGTGTGTGCC
ACGTTGGAGAAGAAGCTCCCGCCCTGCCCCTGACACTGGCCCTTGGGGCCTTCCTGAATCAC

947/6881
FIGURE 881

AAGAACTGGCCTGTACATTTTCAAGGAATTCTTGAGAGGTTCTTGGAGAGATTCTGGGAGCCAAACACTCCATTG
GGATCCTAGCTGTTTTAGAGAACAACCTTGTAATGGAGCCTTCATCTCTTGAGCTGCCGGCTGACACAGTGCAGCG
CATTGCGGCTGAACTCAAATGCCACCCACGGATGAGAGGGTGGCTCTCCACCTAGATGAGGAAGATAAGCTGAG
GCACCTCAGGGAGTGCTTTTATATTCCCAAAATACAGGATCTGCCTCCAGTTGATTTATCATTAGTGAATAAAGA
TGAAAATGCCATCTATTCTTGGGAAATTCTCTTGGCCTTCAACCAAAAATGGTTAAAACATATCTTGAAGAAGA
ACTAGATAAGTGGGCCAAAATAGCAGCCTATGGTCATGAAGTGGGGAAGCGTCCTTGGATTACAGGAGATGAGAG
TATTGTAGGCCTTATGAAGGACATTGTAGGAGCCAATGAGAAAGAAATAGCCCTAATGAATGCTTTGACTGTAAA
TTTACATCTTCTAATGTTATCATTTTTTTAAGCCTACGCCAAAACGATATAAAATTCTTCTAGAAGCCAAAGCCTT
CCCTTCTGATCATTATGCTATTGAGTCACAACCTACAACTTCACGGACTTAACATTGAAGAAAGTATGCGGATGAT
AAAGCCAAGAGAGGGGGAAGAAACCTTAAGAATAGAGGATATCCTTGAAGTAATTGAGAAGGAAGGAGACTCAAT
TGCAGTGATCCTGTTTCACTGGGGTGCAATTTTACACTGGACAGCACTTTAATATTCCTGCCATCACAAAAGCTGG
ACAAGCGAAGGGTTGTTATGTTGGCTTTGATCTAGCACATGCAGTTGGAAATGTTGAACTCTACTTACATGACTG
GGGAGTTGATTTTGCTGCTGGTGTTCCTACAAGTATTTAAATGCAGGAGCAGGAGGAATTGCTGGTGCCTTCAT
TCATGAAAAGCATGCCCATACGATTAAACCTGCATTAGTGGGATGGTTTGGCCATGAACTCAGCACCAGATTTAA
GATGGATAACAACTGCAGTTAATCCCTGGGGTCTGTGGATTCCGAATTTCAAATCCTCCCATTTTGTGGTCTG
TTCCTTGATGCTAGTTTAGAGATCTTTAAGCAAGCGACAATGAAGGCATTGCGGAAAAATCTGTTTGTCTAAC
TGGCTATCTGGAATACCTGATCAAGCATAACTATGGCAAAGATAAAGCAGCAACCAAGAAACCAGTTGTGAACAT
AATTACTCCGTCTCATGTAGAGGAGCGGGGTGCCAGCTAACAATAACATTTTCTGTTCCAAACAAGATGTTTT
CCAAGAACTAGAAAAAAGAGGAGTGGTTTGTGACAAGCGGAATCCAAATGGCATTGAGTGGCTCCAGTTCTCTCT
CTATAATTCTTCCATGATGTTTATAAATTTACCAATCTGCTCACTTCTATACTTGACTCTGCAGAAACAAAAA
TTAGCAGTGTTTTCTAGAACAACCTTAAGCAAATTATACTGAAAGCTGCTGTGGTTATTTAGTATTATTTCGATTT
TTAATTATTGAAAGTATGTCACCATTGACCACATGTAACATAACAATAAATAATATACCTTAC

'948/6881
FIGURE 882

MEPSSLELPADTVQRIAAELKCHPTDERVALHLDEEDKLRHFRECFYIPKIQDLPPVDLSLVNKDENAIYFLGNS
LGLQPKMVKTYLEEEELDKWAKIAAYGHEVGKRPWITGDESIVGLMKDIVGANEKEIALMNALTVNLHLLMLSFFK
PTPKRYKILLEAKAFPSDHYAIESQLQLHGLNIEESMRMIKPREGEETLRIEDILEVIEKEGDSIAVILFSGVHF
YTGQHFNIPAITKAGQAKGCYVGFDLAHAVGNVELYLHDWGVDFAWCSYKYLNAGAGGIAGAFIHEKHAHTIKP
ALVGWFGHELSTRFKMDNKLQLIPGVCGRISNPPILLVCSLHASLEIFKQATMKALRKKSULLTGYLEYLIKHN
YGKDKAATKKPVVNIITPSHVEERGCLTITFSVPNKDVFQELEKRGVVCDKRNPNGIRVAPVPLYNSFHDVYKF
TNLLTSILDSAETKN

949/6881
FIGURE 883

GTGTACGGTCCGCAGCGGCAGGTGAAGTCTAGCAGAGGACGCGGCCAGGCGATTCCGGTGAAGCGATTCTGCAGG
CGTTGGTTCCCTCTTTGACCTGGTAAATGCAGGCTTTTATGCGAGAGGATTTGAATTTGTTGAAATGAGCAGTC
GTAAATCAAAGAGTAACAGCTTAATTCACACAGAGTGCCTTTCACAGGTACAAAGAATTTTACGTGAAAGATTTT
GTCGTGAGAGTCCACATAGTAACCTATTTGGAGTGAAGTACAATACAAACACTTAAGTGAGCTGCTGAAAAGAA
CTGCTCTCCATGGAGAGAGTAACCTCTGTCTTATTATCGGACCCCGAGGATCAGGAAAACTATGTTAATAAATC
ATGCTTTGAAAGAACTCATGGAAATAGAAGAAGTGAGTGAAATGTATTACAAGTTCACCTTAAATGGACTGCTGC
AGATCAATGACAAAATCGCCCTAAAGGAAATCACAAAGGCAGTTAAATCTGGAAAATGTAGTTGGAGATAAAGTTT
TTGGAAGCTTTGCTGAAAACCTTTTCAATTTCTTCTGGAAGCTTTAAAAAAGGTGACCGAACTAGCAGTTGCCAG
TGATCTTCATATTAGATGAATTTGATCTTTTCTGCTCATCATAAAAACCAAACTTCTCTATAATCTTTTGGACA
TTTCTCAGTCTGCACAGACCCCAATAGCAGTTATTGGTCTTACATGTAGATTGGATATTTTGGAACTCTTAGAAA
AAAGAGTGAAGTCAAGATTTTCTCACCAGGCAGATACACTTAATGAATTCATTTGGTTTTCCACAGTATGTTAAAA
TATTTAAAGAACAGTTATCTCTACCTGCAGAGTTTCCAGACAAGGTTTTTGTGAGAAGTGAATGAAAATGTTT
AGTATCTCTCAGAAGATAGAAGTGTGCAAGAAGTACTACAGAAGCATTCAATATCAGCAAAAACCTGCGGTCAT
TACACATGCTATTGATGCTTTGCTTTAAATCGAGTAACAGCATCGCACCCATTTATGACTGCCGTAGATCTAATGG
AAGCAAGCCAACCTGTGTAGCATGGACTCGAAAGCAAATATTGTACATGGTCTATCAGTCTTGAAAATCTGTCTTA
TAATAGCAATGAAACATTAAATGACATCTATGAGGAAGAGCCATTTAATTTTCAAATGGTCTATAATGAGTTTC
AGAAGTTTGTTCAAAAGGAAAGCACATTCGGTTTATAATTTTGAAAAACCTGTGTGTCATGAAGGCTTTTGAACACT
TGCAGCAATTAGAATTAATAAAGCCCATGGAAAGAACTTCAGGAAATTCACAGAGAGAGTACCAGCTGATGAAAC
TGCTTTTGGATAATACTCAAATTATGAATGCTCTGCAGAAATATCCCACTGTCTACAGATGTGAGGCAGTGGG
CAACATCCTCACTAAGCTGGTTATGAATATAACCAGTGACTTCAACTTTGGCATTTCATTCACTTCTGTAGAG
AACGAAAACTATTGTCCATTAACATGATATGCTAAACATTCTATAAACATTCTTGATTTATGTGAGACTTGCC
CATCTACTGTCTTGGCTGTGCTTTGCTTTTAAATCATGAACAGTTACATGATTATAATTTCACTGATTGAGATT
ACTTTGTAAGTAGCTGTTTCAAGAATAAAATATGACTGTTTTAGGGACTAGACCATGTGCTTTTTTAACACTTA
TATATATAATGGTCTATTTGAAGAGCTCACTTCAACCTAACAGCTAGATGTCTTTACAAACCTTAAACCAAAGGA
GTAAAAAACAATGGTAAGCACTGAAGTATAATAAGTAACCTTTGGTACAGCAGGTTTGTGTCAGTGTTTTTTT
CTGTCCACATGCAAATTTTGGATTCTATCCCAGACCCAGGTTTTCTAGTTCAGAAGACTAACAGCTTAGTTCAGA
AGATGGTTCCATGGAAGAAAAAGGCCAAGGAGTTTGAAGATTTTCTTCTAGACATCTCAAAATGTGGTACTCAT
ACCATCCACATCAGAATCCCTTGTAGGATTTTCTAAAATTACAGATTGTTGAGCCTACCATAGGTCAAAAGGACT
GGAATTTTCTTCTTAACAAGTATAGTCATGGCACCAAGTAACTTTTGGTCAATGACATTGTATAAAGGGTGGT
CTCATAAGATTATACCATATTTTACTGTACCTTTTCTATGTCTAAATATACAAATGTTTTACCATTG

950/6881
FIGURE 884

MSSRKSKSNSLIHTECLSQVQRILRERFCRQSPHSNLFQVQVQYKHLSELLKRTALHGESNSVLIIGPRGSGKTM
LINHALKELMEIEEVSENVLQVHLNGLLQINDKIALKEITRQLNLENVVGDKVFGSFAENLSFLLLEALKKGDRTS
SCPVIFILDEFDLFAHHKNQTLTYNLFDISQSAQTPIAVIGLTCRLDILELLEKRVKSRFSHRQIHLMNSFGFPQ
YVKIFKEQLSLPAEFPDKVFAEKWNENVQYLSEDRSVQEVQKHFNISKNLRLHMLLMLALNRVTASHPFMTAV
DLMEASQLCSMDSKANIVHGLSVLEICLI IAMKHLNDIYEEEPFNFMVYNEFQKFVQRKAHSVYNFEKPVVMKA
FEHLQQLELIKPMERTSGNSQREYQLMKLLLDNTQIMNALQKYPNCPTDVRQWATSSLSWL

951/6881
FIGURE 885

ATTCGGCACCGCAGCGTAGGTGCTACCACCGCTGCCGTCGCCGCCCATTTTGATGGCAGGAAGAGTCCGGTTC
TGGGACAGCTGGAGACAGTGGTGGTGAAGTAACTTTACCAAAGGAAAGCTATTTTGCAGAACTATCTTCTCC
AGCGGAGATGGCCAATGTGCTTTGTAAACAGAGCCAGACTGGTTTCCTATCTCCCAGGATTTTGCTCTTTAGTTAA
AAGGGTTGTCAATCCCAAAGCCTTTTCGACTGCAGGATCATCAGGTTTCGGATGAGTCTCATGTGGCTGCTGCACC
TCCAGATATATGCTCTCGAACAGTGTGGCCTGATGAAACTATGGGACCCTTTGGACCTCAAGATCAGAGGTTCCA
GCTTCCTGGGAACATAGGTTTTGATTGTACCTCAATGGGACTGCTTCACAGAAGAAAAGCCTGGTTTCATAAAAC
TTTGCTGATGTTCTAGCAGAACCTTTATCAAGTGAAAGACATGAGTTTGTGATGGCACAATATGTGAATGAATT
TCAGGGTAATGATGCACCTGTTGAACAAGAAATTAACAGTGCAGAAACTTACTTTGAAAGTGCCAGAGTAGAGTG
TGCAATACAAACATGTCCAGAATTGCTGCGAAAAGATTTTGAATCACTGTTTCCAGAAGTAGCTAATGGCAAAC
AATGATTCTGACTGTAAACAAAAAACTAAGAATGATATGACTGTTTGGAGTGAAGAAGTAGAAATTGAAAGAGA
AGTGCTCTTAGAAAAGTTCATCAATGGTGCTAAGGAAATTTGCTATGCTCTTCGAGCTGAGGGTTATTGGGCTGA
CTTTATTGACCCATCATCTGGTTTGGCATTTTTTGGACCATATACAAACAACACTCTTTTTGAAACTGATGAACG
CTACCGACATTTAGGATTCTCTGTTGATGACCTTGGATGCTGTAAAGTGATTTCGTATAGTCTCTGGGGTACCCA
TGTAAGTTGTAGGGAGTATCTTCACTAATGCAACACCAGACAGCCATATTATGAAGAAATTAAGTGGAAATTAGCA
GAAATATCCATTCATTTGCTGTACTATTTGTATGTAATTTTGGGTTGATCTATAAAACACTGTCAGACTAAAGTT
TTTAAATATACTTATTTCTAAGTATTTATTTTCAGCATTTATGAATTTGCAACATTGGCAAGTGATTGGGATTT
TAAATTTGCAATGTTCAATTTATTCATATCATTGAATACACGTTGAACACATCCACATTGTATAGGATGTGGTAA
TTAGCTTGTAACCAGGGTATGATCTGCTATTGTTATTTCTCTCTTTATTGGAAAAAGGCCTCAGTTTTAATTAT
TTTCTTCCCAAAATAAATCACACATTTGGTTACAAAAA

952/6881
FIGURE 886

MANVLCNRARLVSYLPGFCSLVKRVVNPKAFSTAGSSGSDESHVAAAPPDICSRTVWPDETMGPFPGPDQRFQLP
GNIGFDCHLNGTASQKKSLVHKTLPDVLAEP LSSERHEFVMAQYVNEFQGNDA PVEQEINSAETYFESARVECAI
QTCPELLRKDFESLFPEVANGKLMILVTQKTNDMTVWSEEVEIEREV LLEKFINGAKEICYALRAEGYWADFI
DPSSGLAFFGPYTNNTLFETDERYRHLGFSVDDLGCCKVIRHSLWGTHVVVGSIFTNATPDSHIMKKLSGN

953/6881
FIGURE 887

TATCGCAAGCAGCAGTCTCTGGTCCCAGCCCACCCCATGGCCCCCTCCCAGTCCCAGCACCACCAGCAGTAATAAC
AACAGTAGCAGCAGTAGCAACTCAGGATGGGATCAGCTCAGCAAAACGAACCTCTATATCCGAGGACTGCCTCCC
CACACCACCGACCAGGACCTGGTGAAGCTCTGTCAACCATATGGGAAAATAGTCTCCACAAAGGCAATTTTGGAT
AAGACAACGAACAAATGCAAAGGTTATGGTTTTGTGCGACTTTGACAGCCCTGCAGCAGCTCAAAAAGCTGTGTCT
GCCCTGAAGGCCAGTGGGGTTCAAGCTCAAATGGCAAAGCAACAGGAACAAGATCCTACCAACCTCTACATTCT
AATTGGCCACTCTCCATGGATGAGCAAGAACTAGAAAATATGCTCAAACCATTGGACAAGTTATTTCTACAAGG
ATACTACGTGATTCCAGTGGTACAAGTCGTGGTGGTGGCTTTGCTAGGATGGAATCAACAGAAAAATGTGAAGCT
GTTATTGGTCATTTTAATGGAAAAATTTATTAAGACACCACCAGGAGTTTCTGCCCCACAGAACCTTTATTGTGT
AAGTTTGTCTGATGGAGGACAGAAAAAGAGACAGAACCCAAACAAATACATCCCTAATGGAAGACCATGGCATAGA
GAAGGAGAGGTGAGACTTGCTGGAATGACACTTACTTACGACCCAACTACAGCTGCTATACAGAACGGATTTTAT
CCTTACCATAACAGTATTGCTACAAACCGAATGATCACTCAAACCTTCTATTACACCCTATATTGCATCTCCTGTA
TCTGCCTACCAGGTGCAAAGTCCTTCGTGGATGCAACCTCAACCATATATTCTACAGCACCCCTGGTGCCGTGTTA
ACTCCCTCAATGGAGCACACCATGTCTACTACAGCCCGCATCAATGATCAGCCCTCTGGCCCAGCAGATGAGTCAT
CTGTCACTAGGCAGCACCGGAACATACATGCCTGCAACGTCAGCTATGCAAGGAGCCTACTTGCCACAGTATGCA
CATATGCAGACGACGCGGTTCCGTGTTGAGGAGGCAAGTGGTCAACAGCAGGTGGCTGTGAGACGTCTAATGAC
CATTCCTCATATACCTTTCAACCTAATAAGTAAGTGTGAGATGTACAGAAAGGTGTTCTTACATGAAGAAGGGTG
TGAAGGCTGAACAATCATGGATTTTTCTGATCAATTGTGCTTTAGGAAATTATTGACAGTTTTGCACAGGTTCTT
GAAAACGTTATTTATAATGAAATCAACTAAAACTATTTTGTCTATAAGTTCTATAAGGTGCATAAAACCCTTAAA
TTCATCTAGTAGCTGTTCCTCCGAACAGGTTTATTTTAGTAAAAAAAAAAAAA

954/6881
FIGURE 888

MAPPSPTTSSNNSSSSSSNSGWDQLSKTNLYIRGLPPHTTDQDLVKLCQPYGKIVSTKAILDKTTNKCKGYGFV
DFDSPAAQKAVSALKASGVQAQMAKQQEQDPTNLYISNLPLSMDEQELNMLKPFQVISTRILRDSSGTSRGV
GFARMESTEKCEAVIGHFNGKFIKTPPGVSAPTEPLLCKFADGGQKKRQNPKNYIPNGRPWHREGEVRLAGMTLT
YDPTTAAIQNGFYPSYSIATNRMITQTSITPYIASPVSAQVQSPSWMQPQPYILQHFGAVLTPSMEHTMSLQP
ASMISPLAQQMSHLSLGSTGTYPATSAMQGAYLPQYAHMQTTAVPVEEASGQQQVAVETSNDHSPYTFQPNK

955/6881
FIGURE 889

AGACTTGGAAACCCCAAAGTGTCCGCGACCCTGCACGGCAGCTCCCTTCCAGCTTCATGGGCAAAGTGTGGAAAC
AGCAGATGTACCCTCAGTACGCCACCTACTATTACCCCCAGTATCTGCAAGCCAAGTTTGGAAAGGCATTTCGGGAA
TACCAAGTGAAAAGGAAGAGTGAAGAACAGAAAAATGTTAGTGGGATGGCAAACACTGTGAACATTGCTGTTTTCT
AGTGGGCCAGAAAAATCAGTCTCTGGTCCCAGCCCACCCCATGGCCCCCTCCAGTCCCAGCACCCACCAGCAGTAA
TAACAACAGTAGCAGCAGTAGCAACTCAGGATGGGATCAGCTCAGCAAAACGAACCTCTATATCCGAGGACTGCC
TCCCCACACCACCGACCAGGACCTGGTGAAGCTCTGTCAACCATATGGGAAAATAGTCTCCACAAAGGCAATTTT
GGATAAGACAACGAACAAATGCAAAGGTTATGGTTTTGTCGACTTTGACAGCCCTGCAGCAGCTCAAAAAGCTGT
GTCTGCCCTGAAGGCCAGTGGGGTTCAAGCTCAAATGGCAAAGCAACAGGAACAAGATCCTACCAACCTCTACAT
TTCTAATTTGCCACTCTCCATGGATGAGCAAGAACTAGAAAAATATGCTCAAACCATTTGGACAAGTTATTTCTAC
AAGGATACTACGTGATTCCAGTGGTACAAGTCGTGGTGTGGCTTTGCTAGGATGGAATCAACAGAAAAATGTGA
AGCTGTTATTGGTCATTTTAATGGAAAATTTATTAAGACACCACCAGGAGTTTCTGCCCCCAGAACCTTTTATT
GTGTAAGTTTGCTGATGGAGGACAGAAAAAGAGACAGAACCCAAACAAATACATCCCTAATGGAAGACCATGGCA
TAGAGAAGGAGAGGTGAGACTTGCTGGAATGACACTTACTTACGACCCAACCTACAGCTGCTATACAGAACGGATT
TTATCCTTCACCATACAGTATTGCTACAAACCGAATGATCACTCAAACCTTCTATTACACCCTATATTGCATCTCC
TGTATCTGCCTACCAGGTGGCAAAGGAAACCAGAGAAAAACAAGTATCGGGGCTCTGCTATCAAGGTGCAAAGTCC
TTCGTGGATGCAACCTCAACCAATATATTCTACAGCACCTTGGTGCCGTGTTAACTCCCTCAATGGAGCACACCAT
GTCCTACAGCCCGCATCAATGATCAGCCCTCTGGCCCAGCAGATGAGTCATCTGTCACTAGGCAGCACCGGAAC
ATACATGCCTGCAACGTCAGCTATGCAAGGAGCCTACTTGCCACAGTATGCACATATGCAGACGACAGCGGTTCC
TGTTGAGGAGGCAAGTGGTCAACAGCAGGTGGCTGTGAGACGTCTAATGACCATTCTCCATATACCTTTCAACC
TAATAAGTAACTGTGAGATGTACAGAAAGGTGTTCTTACATGAAGAAGGGTGTGAAGGCTGAACAATCATGGATT
TTTCTGATCAATTGTGCTTAGGAAATTATTGACAGTTTTGCACAGGTTCTTGAAAACGTTATTTATAATGAAAT
CAACTAAAACATTTTTTGCTATAAGTTCTATAAGGTGCATAAAACCCTTAAATTCATCTAGTAGCTGTTCCCCCG
AACAGGTTTATTTTAGTAAAAA

956/6881
FIGURE 890

MAPPSPSTTSSNNSSSSSSNSGWDQLSKTNLYIRGLPPHTTDQDLVKLCQPYGKIVSTKAILDKTTNKCKGYGFV
DFDSPAAQKAVSALKASGVQAQMAKQQEQDPTNLYISNLPLSMDEQELENMLKPFQVISTRILRDSSGTSRGV
GFARMESTEKCEAVIGHFNGKFIKTPPGVSAPTEPLLCKFADGGQKKRQNPKNYIPNGRPWHREGEVRLAGMTLT
YDPTTAAIQNGFYSPYSIATNRMITQTSITPYIASPVSAQVAKETRENKYRGSIAKVQSPSWMQPPYILQHP
GAVLTPSMEHTMSLQFASMSPLAQQMSHLSLGSTGYMPATSAMQGAYLPQYAHMQTTAVPVEEASGQQQVAVE
TSNDHSPYTFQPNK

957/6881
FIGURE 891

GGCTAATGTAACATACTCTACCACTTGGTCTGAAGCCCAGCAGTATCTGATGGATAATCCAACCTTTTGCAGAAGA
TGAGGAGTTACAAAATATGGACAAAGAAGATGCATTAATTTGCTTTGAAGAACACATTTCGGGCTTTAGAAAAGGA
GGAAGAAGAAGAAAAACAGAAGAGTTTGTCTGAGAGAAAGGAGACGACAGCGAAAAAATAGGGAATCTTTCCAGAT
ATTTTATAGATGAATTACATGAACATGGACAACCTGCATTCTATGTCATCTTGGATGGAATTGTATCCAACCTATTAG
TTCTGATATTAGATTCACTAATATGCTTGGTCAGCCTGGATCAACTGCACCTTGATCTTTTCAAGTTTTATGTTGA
GGATCTTAAAGCACGTTATCATGACGAGAAGAAGATAATAAAAGACATTCTAAAGGATAAAGGATTTGTAGTTGA
AGTAAACACTACTTTTGAAGATTTTGTGGCGATAATCAGTTCAACTAAAAGATCAACTACATTAGATGCTGGAAA
TATCAAATTGGCTTTCAATATTACTAGAAAAGGCAGAAGCCCGTGAACGTGAAAGAGAAAAAGAGAGGCTCGGA
AGATGAAACGAAAAGAATCTGCATTTAAGAGTATGTTAAAAACAAGCTGCTCCTCCGATAGAATTGGATGCTGTCT
GGGAAGATATCCGTGAGAGATTTGTAAAAGAGCCAGCATTGAGGACATAACTCTAGAATCTGAAAGAAAACGAA
TATTTAAAGATTTTATGTCATGTGCTTGAGCATGAATGTCAGCATCATCATTCAAAGAACAAGAAACATTCTAAGA
AATCTAAAAAACATCATAGGAAACGTTCCCGCTCTCGATCGGGGTCAGATTGAGATGATGATGATAGCCATTCAA
AGAAAAAAGACAGCGATCAGAGTCTCGTTCTGCTTCAGAACATTCTTCTAGTGCAGAGTCTGAGAGAAGTTATA
AAAAGTCAAAAAAGCATAAGAAGAAAAGTAAGAAGAGGAGACATAAATCTGACTCTCCAGAATCCGATGCTGAGC
GAGAGAAGGATAAAAAAGAAAAGATCGGGAAAGTGAAAAAGACAGAACTAGACAAAGATCAGAATCAAAACACA
AATCGCCTAAGAAAAAGACTGGAAGGATTCTGGTAATTGGGATACTTCTGGCAGCGAACTGAGTGAAGGGGAAT
TGGAAAAGCGCAGAAGAACCCTTTTGGAGCAACTGGATGATGATCAATAAATTATACCAAATATATGTTTACAGT
ATGATTTAAAGTCTGATTCAGACCAGGGACTCTATTTTAAAGTTCAACTGAAATAACACTGGGTTTTAATTATATC
ACAGGAAAAAAAAGTGCATTTAAGTATTGTTATCGTGGACTTTATAAAAGCAAAGGAAATTGAAAGTAACTTTT
GATTCTGTATCAAGAATCATATTTTCATACAGTCATAACTGTCTTTCTGTGACCCTTTACAGGGCACTGTAGGA
TGGATTAAAGGTGGCAATTTACTGATAACTGCAGATGTCTCTACTTTGTTCTAAAAATCTAAGTCATGAGGTGATT
TGATTTACTTTATAGAAGCTGGATTTTGAAGATCTAATGAAAAATTTTTTGATAATATAGTAGTACAAAAAAGC
ACCAGCAACTGATAAAAAATTGCTTTTTTGTGCGCTACCCAACCTGGTTAAAGCCAATGTGATCTTTTATGGTGAAA
CTCCTAAGAAACAGGTGGTTTTGCTGGAACTTGGTAGACCCTTAATTATAGTGGTGCTAATGAGCACTACTGTA
ATATAAAGCCACCATTATTTTTATCAAACATCTGAATACATTTTACAAAGGCTATTGTGAGGGCATTATTTTGA
GCATCTATTTGAGGTGATGTTAAAAAACTTTAACATCAAATCAAATTGTAATTAATTTAAATATATTGCCT
TAAGGACCTACTAAAGAATGTGCCACCAGACTTTAAGTGATAGTTGCAATATCCTTGCTCTAAAAAAGGCTGCTG
GTTGACTTAAACATTTTCTTTAACAGTTGTCTTTTTTTCTAAATTCAGTCTTCTCTTGCTTTTTTTTCCCTGC
TATTGAGGAAGTATTTTGCCTTCCCTACTCACTGAGAAGTATTGACTTCGTGGTACACATTCTAAAGCATTCTG
ATTTGAATATTTTTGTACATTTTATCAATTATTAACCTTCTCTCTAGTG

958/6881
FIGURE 892

MLEISNWLSILLEKAEAREREREKEEARKMKRKESAFKSMKQAAPPIELDAVWEDIRERFVKEPAFEDITLESE
RKRIKDFMHVLEHECQHHSKNKKHSKSKKHHRKRSRSGSDSDDDSHSKKRQRSESRASEHSSSAESE
RSYKSKKHKKSKRRHKSDSPESDAEREKDKKEKDRESEKDRTRQRSESKHKSPKKKTGKDSGNWDTSGSELS
EGELEKRRRTLLEQLDDQ

959/6881
FIGURE 893

CCAGCACGCGGCCGCTCTGCCCCAGGGAAGTCCCTGTCTTACCTTCAGCAGGAGCCGGTTCCCTGTGTGTGTGT
CCGCTCGCCCTCTGCTCCGTCTGCGGCTGCCCCACTGCCCTCTACGGTCCACCATGGCCCTGCTGCACTCCGGC
CGCGTCTCTCCCGGGATCGCCGCCGCTTCCACCCGGGCTCGCCGCCGCGGCTCTGCCAGAGCCAGCTCTGG
TGGACCCATGTGGAAATGGGACCTCCAGATCCATTCTGGGAGTCACTGAAGCCTTTAAGAGGGACACCAATAGC
AAAAAGATGAATCTGGGAGTTGGTGCTACCGGGATGATAACGGAAAGCCTTACGTGCTGCCTAGCGTCCGCAAG
GCAGAGGCCAGATTGCCGCAAAAAATTGGACAAGGAATACCTGCCCATTTGGGGGACTGGCTGAATTTTGCAAG
GCATCTGCAGAACTAGCCCTGGGTGAGAACAGCGAAGTCTTGAAGAGTGGCCGGTTTGTCACTGTGCAGACCATT
TCTGGAAGTGGAGCCTTAAGGATCGGAGCCAGTTTTCTGCAAGATTTTTTAAGTTTCAGCCGAGATGTCTTTCTG
CCCAAACCAACCTGGGGAAACCACACACCCATCTTCAGGGATGCTGGCATGCAGCTACAAGGTTATCGGTATTAT
GACCCCAAGACTTGGGTTTTGACTTCACAGGCGCTGTGGAGGATATTTCAAAAATACCAGAGCAGAGTGTTCTT
CTTCTGCATGCCTGCGCCACAATCCCACGGGAGTGGACCCGCGTCCGGAACAGTGAAGGAAATAGCAACAGTG
GTGAAGAAAAGGAATCTCTTTGCGTTCTTTGACATGGCTACCAAGGCTTTGCCAGTGGTGATGGTGATAAGGAT
GCCTGGGCTGTGCGCCACTTCATCGAACAGGGCATTAAATGTTTGTCTCTGCCAATCATATGCCAAGAACATGGGC
TTATATGGTGAGCGTGTAGGAGCCTTCACTATGGTCTGCAAAGATGCGGATGAAGCCAAAAGGGTAGAGTCACAG
TTGAAGATCTTGATCCGTCCCATGTATTCCAACCCCTCCCTCAATGGGGCCCGGATTGCTGCTGCCATTCTGAAC
ACCCAGATTTGCGAAAACAATGGCTGCAAGAAGTGAAGGCGATGGCTGACCGCATCATTTGGCATGCGGACTCAA
CTGGTCTCCAACCTCAAGAAGGAGGGTTCCACCCACAATTGGCAACACATCACCGACCAAATTGGCATGTTCTGT
TTCACAGGGCTAAAGCCTGAACAGGTGGAGCGGCTGATCAAGGAGTTCTCCATCTACATGACAAAAGATGGCCGC
ATCTCTGTGGCAGGGGTACCTCCAGCAACGTGGGCTACCTTGCCCATGCCATTACCAGGTCACCAAGTAATGT
CCCTGGTGCGAGGAAACAGAGACAACCTTTCTGTCTTCAGCCTCTGCTATTGAGAGCTTCACACAGACAATGAGA
GAGGGTGGATGGTGGTGAGTGGATCATTCTTTTCAGCCACAGTGTGTAACACTCAGCATTGGAATGTTTCTCAGA
AAAGAACATGTAGTGACACAGGGCAGAGGCATCCATGGCTGGCGTCTGGAATATTAACCAAACCTCTCCCGGTC
CTTTTTCTCCAACCTTTCTCAAAGAGTTTACATGTGCAAGAAAGTCATCGCACCAAAAAACCTGTCAATTATGC
CATTGCAATATTTTCAGAAGCTTTAACTGAAGTGTGAGGTTCCCTCGTGAGAAACAGCACACGTTAGAGGCTTTGAG
AGAAGGCCTAGTTCTGTATGATGATGTCGCGCTCGTGTCTGTCCTCCCATCTTGGAACAACCTTATCAACAGGCC
GCACTGCAGAAATGATGTTTTATGAAAACCAATGAGGCTGCTGCCACTCCAGCAAGGGAAATAATGCAGTTTCCT
GTCTTATTTAAGAAAAAGAGAAGGCTCTCTTTCTCCCTTGTCATTGCCGTTCTTTTCTTACACGCAAAGATTT
TTAACTATTGCAGATTTTCATCCATTCTACTGCTTGATTGACCATCAACTCCATCCTATCGAGATTTATTTAA
GAATGAAGAACATAATTTCTGCTGATGCCGTACCTCACCTTTTCAGCAAAGAATAGTGGAGAGTAGGAACT
GTACTTTATCTCGGCATCCTCTTGAATGATAGTGCAAGTTTCTCCAGTTGGGATGTTGTCTCTGCCCGGTTGGAC
CTCCTCCCTTTGTTGAATGTGGTGTGCAGCCTCTCATCTCACACTGTGAGTCCAGCGGCGCAGGGTGGTACCAGG
AAAGAGGATATTCTAGGCTTTGCGTGCTGCTAGCTGGGTTCAGGCTTCACCCACTGGAAAGAACCACCATCTGCT
CTAACCATGTAGACTTATTGCGGCCTGGTTTTCTGTGTACAATAAAATTACTGTAGACC

960/6881
FIGURE 894

GAGTGACTCCACCGCCCGGAGCAGCGGTGCAGGACGCGCGTCTCCGCCGCCCGGGTGACTTCTGCCTGCGCTCC
TTCTCTGAACGCTCACTTCCGAGGAGACGCCGACGATGAAGACACCGTGGAAGGTTCTTCTGGGACTGCTGGGTG
CTGCTGCGCTTGTCAACATCATCACCGTGCCCGTGGTTCTGCTGAACAAAGGCACAGATGATGCTACAGCTGACA
GTGCAAAACTTACACTCTAACTGATTACTTAAAAATACTTATAGACTGAAGTTATACTCCTTAAGATGGATTT
CAGATCATGAATATCTCTACAAACAAGAAAATAATATCTTGGTATTCAATGCTGAATATGGAAACAGCTCAGTTT
TCTTGAGAACAGTACATTTGATGAGTTGGACATTCTATCAATGATTATTCAATATCTCCTGATGGGCAGTTTA
TTCTCTTAGAATACAACCTACGTGAAGCAATGGAGGCATTCTACACAGCTTCATATGACATTTATGATTTAAATA
AAAGGCAGCTGATTACAGAAGAGAGGATTCCAAACAACACAGTGGGTCACATGGTCACCAGTGGGTCATAAAT
TGGCATATGTTTGGAACAATGACATTTATGTTAAATTTGAACCAAATTTACCAAGTTACAGAATCACATGGACGG
GGAAAGAAGATATAATATATAATGGAATAACTGACTGGGTTTATGAAGAGGAAGTCTTCAGTGCCCTACTCTGCTC
TGTGGTGGTCTCCAAACGGCACATTTTTTAGCATATGCCCAATTTAACGACACAGAAGTCCCACTTATTGAATACT
CCTTCTACTCTGATGAGTCACTGCAGTACCCAAAGACTGTACGGGTTCCATATCCAAAGGCAGGAGCTGTGAATC
CAACTGTAAAGTTCTTTGTTGTAATACAGACTCTCTCAGCTCAGTCACCAATGCAACTTCCATACAAATCACTG
CTCCTGCTTCTATGTTGATAGGGGATCACTACTTGTGTGATGTGACATGGGCAACACAAGAAAGAAATTTCTTTGC
AGTGGCTCAGGAGGATTCAGAATATTCCGGTCATGGATATTGTGACTATGATGAATCCAGTGGGAAGATGGAAT
GCTTAGTGGCAGGCAACACATTGAAATGAGTACTACTGGCTGGGTTGGAAGATTTAGGCCTTCAGAACCTCATT
TTACCCCTTGATGGTAATAGCTTCTACAAGATCATCAGCAATGAAGAAGGTTACAGACACATTTGCTATTTCCAAA
TAGATAAAAAAGACTGCACATTTATTACAAAAGGCACCTGGGAAGTCATCGGGATAGAAGCTCTAACCAAGTGATT
ATCTATACTACATTAGTAAATGAATATAAAGGAATGCCAGGAGGAAGGAATCTTTATAAAATCCAACCTTAGTGACT
ATACAAAAGTGACATGCCCTCAGTTGTGAGCTGAATCCGGAAGGTGTGAGTACTATTCTGTGTCATTTCAGTAAAG
AGGCGAAGTATTATCAGCTGAGATGTTCCGGTCTCGGCTCTATCTCTACACAGCAGCGTGAATGATA
AAGGGCTGAGAGTCTGGAAGACAATTCAGCTTTGGATAAAATGCTGCAGAAATGTCAGATGCCCTCCAAAAAAC
TGGACTTCATTATTTTGAATGAAACAAAATTTTGGTATCAGATGATCTTGCCTCCTCATTTTGATAAATCCAAGA
AATATCCTCTACTATTAGATGTGTATGCAGGCCCATGTAGTCAAAAAGCAGACACTGTCTTCAGACTGAACTGGG
CCACTTACCTTGCAAGCACAGAAAACATTATAGTAGCTAGCTTTGATGGCAGAGGAAGTGGTTACCAAGGAGATA
AGATCATGCATGCAATCAACAGAAGACTGGGAACATTTGAAGTTGAAGATCAAATTGAAGCAGCCAGACAATTTT
CAAAAATGGGATTTGTGGACAACAAACGAATTTGCAATTTGGGGCTGGTCATATGGAGGGTACGTAACTCAATGG
TCCTGGGATCGGGAAGTGGCGTGTCAAGTGTGGAATAGCCGTGGCGCCTGTATCCCGGTGGGAGTACTATGACT
CAGTGTACACAGAACGTTACATGGGTCTCCCAACTCCAGAAGACAACCTTGACCATTACAGAAATTCACAGTCA
TGAGCAGAGCTGAAAATTTTAAACAAGTTGAGTACCTCCTTATTCATGGAACAGCAGATGATAACGTTCACTTTC
AGCAGTCAGCTCAGATCTCCAAAGCCCTGGTCGATGTTGGAGTGGATTTCAGGCAATGTGGTATACTGATGAAG
ACCATGGAATAGCTAGCAGCACAGCACACCAACATATATATACCCACATGAGCCACTTCATAAAACAATGTTTCT
CTTTACCTTAGCACCTCAAAATACCATGCCATTTAAAGCTTATTTAAACTCATTTTTGTGTTTCATTATCTCAAAA
CTGCACTGTCAAGATGATGATGATCTTTAAATACACACTCAAATCAAGAACTTAAGGTTACCTTTGTTCCCAA
ATTTACATACCTATCATCTTAAGTAGGGACTTCTGTCTTCACAACAGATTATTACCTTACAGAAGTTTGAATTATC
CGGTGCGGTTTTATTGTTTAAATCATTCTGTCATCAGCTGCTGAAACAACAAATAGGAATTGTTTTTATGGAGG
CTTTGCATAGATTCCCTGAGCAGGATTTTAACTTTTTCTAACTGGACTGGTTCAAATGTTGTTCTCTTTTAA
AGGGATGGCAAGATGTGGCAGTGATGTCACTAGGGCAGGGACAGGATAAGAGGGATTAGGGAGAGAAGATAGCA
GGGCATGGCTGGGAACCAAGTCCAAGCATAACACAGCAGGCTACTGTGAGCTCCCTCGGAGAAGAGCTG
TTCACAGCCAGACTGGCACAGTTTCTGAGAAAGACTATTCAAACAGTCTCAGGAAATCAAATATGCAAAGCACT
GACTTCTAAGTAAAACACAGCAGTTGAAAAGACTCCAAAGAAATGTAAGGGAACTGCCAGCAACGCAGGCCCC
CAGGTGCCAGTTATGGCTATAGGTGCTACAAAACACAGCAAGGGTGATGGGAAAGCATTGTAAATGTGCTTTTA
AAAAAAATACTGATGTTCCTAGTGAAAGAGGCAGCTTGAAACTGAGATGTGAACACATCAGCTTGCCCTGTTAA
AAGATGAAAATATTTGTATCACAATCTTAACCTGAAGGAGTCCTTGCAATCAATTTTCTTATTTCATTCTTTG
AGTGTCTTAATTAAGAATATTTTAACTTCCTTGAGTCAATTTTAAAAATGGAACATAAAATACAATGTTATG
TATTATTATCCCACTTCTACATACTATGGAATTTCTCCAGTCATTTAATAAATGTGCCTTCATTTTTTC

961/6881
FIGURE 895

MKTPWKVLLGLLGAAALVTIITVPVLLNKGTDATADSRKTYTLTDYLKNTYRLKLYSLRWISDHEYLYKQENN
ILVFNAEYGNSSVFLENSTFDEFGHSINDYSISPDGQFILLEYNVVKQWRHSYTASYDIYDLNKRQLITEERIPN
NTQWVTWSPVGHKLAYVWNNDIYVKIEPNLPSYRITWTGKEDIYNGITDWVYEEEVFSAYSALWWSPNGTFLAY
AQFNDTEVPLIEYSFYSDSLQYPKTVRVPYPKAGAVNPTVKFFVVNTDSLSSVTNATSIQITAPASMLIGDHYL
CDVTWATQERISLQWLRRIQNYSVMDICDYDESSGRWNCVARQHIEMSTTGWVGFRFPSEPHFTLDGNSFYKII
SNEEGYRHICYFQIDKKDCTFITKGTWEVIGIEALTSYLYYISNEYKGMPPGGRNLYKIQLSDYTKVTCLSCELN
PERCQYYSVSFSKEAKYYQLRCSGPGPLPLYTLHSSVNDKGLRVLEDNSALDKMLQNVQMPSKKLDFIILNETKFW
YQMILPPHFDKSKKYPLLLDVYAGPCSQKADTVFRLNWATYLASTENIIVASFDRGSGYQGDKIMHAINRRLGT
FEVEDQIEAARQFSKMGFVDNKRIAIWGWSYGGYVTSMVLGSGSGVFKCGIAVAPVSRWEYDYSVYTERYMGLPT
PEDNLDHYRNSTVMSRAENFKQVEYLLIHGTADDNVHFQQSAQISKALVDVGVDFOAMWYTDEDHGIASSTAHQH
IYTHMSHFQCFSLP

962/6881
FIGURE 896

GT TTGAAATCGGAAAGTTGGCGGGGCTGCGGGAGCTGAGCCTAGAGTCCGGCTGTTGGCTAGAGTGGGCGCGGAT
CTGGTGTGGGGAAGGCGGCGGGACTCAGGCCTGCCTGCÁAGCATTGTCTACATAATGGTAGAGGACGAACTGG
CACTTTTCGATAAAAGCATAAATGAATTTTGAATAAATTCAAAGTACGGACACCTCCTGTCAGATGGCGGGAC
TAAGAGATACCTACAAGGATTCCATCAAAGCATTTCAGAAAAGCTGTCTGTGAAATTAAGGAAGAAGAACC
TGGTTGAGATGTTTCTGGAATATCAAATCAGATCAGCAGGCAAAATAAGCTCATTCAAGAAAAAAGGATAACT
TGTTAAATTTGATTGCTGAAGTAAAGGCAAAAGCAGGAATTGGAAGTACTGACTGCAATATCCAGGATCTTA
AGGAAGAATATTCTAGGAAGAAGGAACTATTTCTACTGCTAATAAGCGAATGCAGAGAGGTTGAAAAGGCTGC
AGAAATCTGCAGACTTGTATAAGATCGACTTGGACTAGAAATTCGAAAAATTTATGGTGAGAAATTGCAGTTTA
TTTTCACTAATATTGACCCTAAGAATCCTGAGAGCCCATTTATGTTTTCTTACATCTCAATGAAGCAAGGGACT
ATGAAGTGTGAGATAGTCCCCCTCATCTTGAGGGCCTAGCAGAATTTCAAGAGAATGTAAGGAAGACCAACAATT
TTTCAGCTTTTCTTGCCAATGTTTCGAAAGCTTTTACTGCCACGGTTTATAATTAACATACAAATAGTGATATA
AAAACGGTTTATTTTCTTCTCTATTACATATCTTTTTTTTTCTTGTTTTATTATTACTATACITTAAGTTTT
AGGGTACATGTGCACAATGTGCAGGTTTGTACATATGTATACATGTGCCATATTGGTGIGCTGCACCCATTAAC
TCGTCAATTCATTAGGTATATCTCCTAATGCTATCCCTCCCCCTCCCCCAACCCACAACAGTCCCCGTTGTGTG
ATGTTCCCTTCCTGTGTCCATGTGTTCTCATTGTTCAATTCACCTAGGAGTGAGAATATGTGGTGTGTTGGTT
TTTTGTCTTTTCGATAGTTTGCTGAGAATGATGGTTTCCAGCTTCATCCATGTTCTTACAAAGGACATGAATCA
TCCTTTTTTATGGCTGCATAGTATTCCATGGTGTATATGTGCCACATTTTCTTAATCCAGTCTATCATTGTTGGA
CATTTGGGTTGGTTCCAAGTCTTTGCTATTGTGAATAGTGCCGAAATAAACATACGTGTGCATGTGTCTCCAAAA
AAAAAAAAAAAAAAAAAAAAAAAAA

963/6881
FIGURE 897

MVEDELALFDKSINEFWNKFKSTDTSCQMAGLRDITYKDSIKAFAEKLSVKLKEEERMVEMFLEYQNQISRQNKLI
QEKKNLLKLI AEVKGKKQELEVLTANIQDLKEEYSRKKETISTANKANAERLKRLOKSADLYKDRGLGLEIRKIY
GEKLQFIFTNIDPKNPESPFMFSLHLNEARDYEVSDSAPHLEGLAEFQENVRKTNNFSAFLANVRKAFTATVYN

964/6881
FIGURE 898

AAGCACAGGCCACCACTCTGCCCTGGTCCACACAAGCTCCGGTAGCCCATGAGGCCCTGGCCTCTCCTCCTGCTC
TTTAGCCCTTTGCTCAGCTGGCCTCGTCCTGGGCTCCGAACATGAGACCCGTCTGGTGGCAAAGCTATTTAAAGAC
TACAGCAGCGTGGTGCGGCCAGTGGAAGACCACCGCCAGGTCGTGGAGGTCACCGTGGGCCTGCAGCTGATACAG
CTCATCAATGTGGATGAAGTAAATCAGATCGTGACAACCAATGTGCGTCTGAAACAGCAATGGGTGGATTACAAC
CTAAAAATGGAATCCAGATGACTATGGCGGTGTGAAAAAATTACATTCCCTTCAGAAAAGATCTGGCGCCCAGAC
CTTGTCTCTATAACAATGCAGATGGTGACTTTGCTATTGTCAAGTTCACCAAAGTGCTCCTGCAGTACACTGGC
CACATCACGTGGACACCTCCAGCCATCTTTAAAGCTACTGTGAGATCATCGTCACCCACTTTCCCTTTGATGAA
CAGAACTGCAGCATGAAGCTGGGCACCTGGACCTACGACGGCTCTGTCTGGCCATCAACCCGGAAAGCGACCAG
CCAGACCTGAGCAACTTCATGGAGAGCGGGGAGTGGGTGATCAAGGAGTCCCGGGGCTGGAAGCACTCCGTGACC
TATTCCTGCTGCCCCGACACCCCTACCTGGACATCACCTACCACTTCGTTCATGCAGCGCCTGCCCTCTACTTC
ATCGTCAACGTCATCATCCCCCTGCCTGCTCTTCTCCTTCTTAACTGGCCTGGTATTCTACCTGCCACAGACTCA
GGGGAGAAGATGACTCTGAGCATCTCTGTCTTACTGTCTTTGACTGTGTTCCTTCTGGTCAICGTGGAGCTGATC
CCCTCCACGTCCAGTGCTGTGCCCTTGATTGGAATAACATGCTGTTCACCATGGTGTTCGTTCATTGCCTCCATC
ATCATCACTGTTCATCGTCATCAACACACACCACCGCTCACCCAGCACCCATGTTCATGCCCAACTGGGTGCGGAAG
GTTTTTATCGACACTATCCCAAATATCATGTTTTTCTCCACAATGAAAAGACCATCCAGAGAAAAGCAAGACAAA
AAGATTTTACAGAAGACATTGATATCTCTGACATTTCTGGAAAGCCAGGGCCTCCACCCATGGGCTTCCACTCT
CCCCTGATCAAACACCCCGAGGTGAAAAGTGCCATCGAGGGCATCAAGTACATCGCAGAGACCATGAAGTCAGAC
CAGGAGTCTAACAATGCGGCGGCAGAGTGGAAAGTACGTTGCAATGGTGATGGACCACATACTCCTCGGAGTCTTC
ATGCTTGTTTGCAICATCGGAACCTAGCCGTGTTTGCAAGTTCGACTCATTGAATTAAATCAGCAAGGATGAGCA
GAAAATGAGCTGAGCTTAGCTCTGCCCTGGAACCTACCAGAGCAGAGAAGGGCAGGAGAGGAAGATTGTCTACT
TGCTCCACTCGCACTTATCAAACGTGTTATATCCATACTTATTATTGATGATAAGATTTACCTTTATGTAAGTT
TATGGCCTTGAAGTGTTTTTCATATTGCTTCTCCCTTTAGTTCTGCTGTCTCCCTGAAGAGTGAACCTCTTTAGT
AAATGAAACTAATCACT

965/6881
FIGURE 899

MEPWPLLLLFSLCSAGLVLGSEHETRLVAKLFKDYSSVVRPVEDHRQVVEVTVGLQLIQLINVDEVNQIVTTNVR
LKQQWVDYNLKNPDDYGGVKKIHIPSEKIWRPDLVLYNNADGDFAIVKFTKVLLQYTGHITWTFFAIFKSYCEI
IVTHFPFDEQNCSMKLGTWYDGSVVAINPESDQPDLSNFMESGEWVIKESRGWKHSVTYSCCPDTPYLDITYHF
VMQRLPLYFIVNVIIPCLLFSFLTGLVFYLPDTSGEKMTLSISVLLSLTVFLLVIVELIPSTSSAVPLIGKYMFL
TMVFVIASIIITVIVINTHHRSPSTHVMPNWRKVFIDTIPNIMFFSTMKRPSREKQDKKIFTEDIDISDISGKP
GPPPMGFHSPLIKHPEVKS AIEGIKYIAETMKSDQESNNAAEWKYVAMVMDHILLGVFMLVCIIGTLAVFAGRL
IELNQOG

966/6881
FIGURE 900

GTTTCGCAGCCGCCGCCGCCGCCGCTCGCTCTCCAACGCCAGCGCCGCCCTCTCGCTCGCCGAGCTCCAGCCGAAG
GAGAAGGGGGGTAAGTAAGGAGGTCTCTGTACCATGGCTCGTACAAAGCAGACTGCCCCGAAATCGACCGGTGGT
AAAGCACCCAGGAAGCAACTGGCTACAAAAGCCGCTCGCAAGAGTGCGCCCTCTACTGGAGGGGTGAAGAAACCT
CATCGTTACAGGCCTGGTACTGTGGCGCTCCGTGAAATTAGACGTTATCAGAAGTCCACTGAACTTCTGATTTCGC
AAACTTCCCTTCCAGCGTCTGGTGCGAGAAATTGCTCAGGACTTTAAAACAGATCTGCGCTTCCAGAGCGCAGCT
ATCGGTGCTTTGCAGGAGGCAAGTGAGGCCTATCTGGTTGGCCTTTTTGAAGACACCAACCTGTGTGCTATCCAT
GCCAAACGTGTAACAATTATGCCAAAAGACATCCAGCTAGCACGCCGCATACGTGGAGAACGTGCTTAAGAATCC
ACTATGATGGGAAACATTTTCTCTCAAAAAAAAAAAAAAAAAATTTCTCTTCTTCTGTTATTGGTAGTTCTGAAC
GTTAGATATTTTTTTTCCATGGGGTCAAAAGGTACCTAAGTATATGATTGCGAGTGGAAAAATAGGGGACAGAAA
TCAGGTATTGGCAGTTTTTCCATTTTCATTTGTGTGTGAATTTTAAATATAAATGCGGAGACGTAAAGCATTAAAT
GCAAGTTAAATGTTTCAGTGAACAAGTTTCAGCGGTTCACTTTATAATAATTATAAATAAACCTGTAAATTT
TTCTGGACAATGCCAGCATTTGGATTTTTTTAAAACAAGTAAATTTCTTATTGATGGCAACTAAATGGTGTGTGT
AGCATTTTTATCATACAGTAGATTCCATCCATTCACTATACTTTTCTAACTGAGTTGTCCTACATGCAAGTACAT
GTTTTTAATGTTGTCTGTCTTCTGTGCTGTTTCTGTAAGTTTGCTATTAAAATACATTAAACTAT

967/6881
FIGURE 901

CGCCGAGCTCCAGCCGAAGGAGAAGGGGGTAAGTAAGGAGGTCTCTGTACCATGGCTCGTACAAAGCAGACTGC
CCGCAAATCGACCGGTGGTAAAGCACCCAGGAAGCAACTGGCTACAAAAGCCGCTCGCAAGAGTGCGCCCTCTAC
TGGAGGGGTGAAGAAACCTCATCGTTACAGGCCTGGTACTGTGGCGCTCCGTGAAATTAGACGTTATCAGAAGTC
CACTGAACTTCTGATTCGAAACTTCCCTTCCAGCGTCTGGTGCGAGAAATGCTCAGGACTTTAAAACAGATCT
GCGCTTCCAGAGCGCAGCTATCGGTGCTTTGCAGGAGGCAAGTGAGGCCTATCTGGTTGGCCTTTTGAAGACAC
CAACCTGTGTGCTATCCATGCCAAACGTGTAACAATTATGCCAAAAGACATCCAGCTAGCACGCCGCATACGTGG
AGAACGTGCTTAAGAATCCACTATGATGGGAAACATTTCACTCTCTCCATGGGGTCAAAAGGTACCTAAGTATAT
GATTGCGAGTGAAA

968/6881

FIGURE 902

MARTKQTARKSTGGKAPRKQLATKAARKSAPSTGGVKKPHRYRPGTVALREIRRYQKSTELLIRKLPFQRLVREI
AQDFKTDLRFQSAAGALQEASEAYLVGLFEDTNLCAIHAKRVTIMPKDIQLARRIRGERA

969/6881
FIGURE 903

CAGCCGTTGAGGGGACGGGCCTGCGTTCTCTCCTCCTTCCCTCCCGCCTCCAGCTGCCGGCAGGACCTTTCTCTC
GCTGCCGCTGGGACCCCGTGTCTATCGCCAGGCCGAGCACGATGCCCCCTAAAAAGGGAGGTGATGGAATTAAAC
CACCCCAATCATTGGAAGATTTGGAACCTCACTGAAAATTGGTATTGTTGGATTGCCAAATGTTGGGAAATCTA
CTTTCTTCAATGIGTTAACCAATAGTCAGGCTTCAGCAGAAAACTTCCCGTTCTGCACTATTGATCCTAATGAGA
GCAGAGTACCTGTGCCAGATGAAAGGTTTGACTTTCTTTGTCAATACCACAAACCAGCAAGCAAAATTCCTGCCT
TTCTAAATGTGGTGGATATTGCTGGCCTTGTGAAAGGAGCTCACAATGGGCAGGGCCTGGGGAATGCTTTTTTAT
CTCATATTAGTGCCTGTGATGGCATCTTTCATCTAACACGTGCTTTTGAAGATGATGATATCACGCACGTTGAAG
GAAGTGTAGATCCTATTTCGAGATATAGAAATAATACATGAAGAGCTTCAGCTTAAAGATGAGGAAATGATTGGGC
CCATTATAGATAAACTAGAAAAGGTGGCTGTGAGAGGAGGAGATAAAAACTAAAACTGAATATGATATAATGT
GCAAAGTAAAATCCTGGGTATAGATCAAAAGAAACCTGTTTCGCTTCTATCATGATTGGAATGACAAAGAGATTG
AAGTGTGAAATAAACACTTATTTTTGACTTCAAAACCAATGGTCTACTTGGTTAATCTTTCTGAAAAAGACTACA
TTAGAAAGAAAAACAAATGGTTGATAAAAAATTAAAGAGTGGGTGGACAAGTATGACCCAGGTGCTTTGGTCATTC
CTTTTAGTGGGGCCTTGGAACCTCAAGTTGCAAGAATTGAGTGTGAGGAGAGACAGAAGTATCTGGAAGCGAACA
TGACACAAAGTGCTTTGCCAAAGATCATTAAAGGCTGGGTTTGCAGCACTCCAACTAGAATACTTTTTCACTGCAG
GCCCAGATGAAGTGCGTGCATGGACCATCAGGAAAGGGACTAAGGCTCCTCAGGCTGCAGGAAAGATTACACAG
ATTTTGAAGAGGATTCATTATGGCTGAAGTAATGAAATACGAAGATTTTAAAGAGGAAGGTTCTGAAAATGCAG
TCAAGGCTGCTGGAAAGTACAGACAACAAGGCAGAAATTATATTGTTGAAGATGGAGATATTATCTTCTTCAAAT
TTAACACACCTCAACAACCGAAGAAGAAATATAATTTAGTTATTGCTCAGATAAACATACAACCTCCAAAAGGCA
TCTGATTTTTTAAAAAATTAAAAATTTCTGAAAACCAATGCGACAAATAAAGTTGGGGAGATGGGAATCTTTGACAA
ACAAATTATTTTTATTTGTTTTTAAATTAATACTGTGTACCCCCCCCCCCCCATGAAATGCAGGTTCACTAAA
TGTGAACAGCTTTGCTTTTCACGTGATTAAAGACCCTACTCCAAATTGTAGAAGCTTTTCAGGAACCATATTACTC
TCATGATACTTCATTAATCTCCATCAIGTATGCCAAGCCTGACACATTTGACAGTGAGGACAATGTGGCTTGCTC
CTTTTGAATCTACAGATAATGCATGTTTTACAGTACTCCAGATGTCTACACTCAATAAAACATTTGACAAAACC
AAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA

970/6881
FIGURE 904

MPPKKGGDGIKPPPIIGRFGTSLKIGIVGLPNVGKSTFFNVLTNSQASAENFPFCTIDPNESRVVPDERFDFLC
QYHKPASKIPAFNLNVVDIAGLVKGAHNGQGLGNAFLSHISACDGIFHLTRAFEDDDITHVEGSVDPIRDIEIHE
ELQLKDEEMIGPIIDKLEKVAVRGGDKKLKPEYDIMCKVKSVIDQKKPVRFYHDWNDKEIEVLNKHFLTSPKM
VYLVNLSEKDYIRKKNKWLIIKIKWVDKYDPGALVIPFSGALELKLQELSAEERQKYLEANMTQSALPKI KAGF
AALQLEYFFTAGPDEVRAWTIRKGTAPQAAGKIHTDFEKGFI MAEVMKYEDFKEEGSENAVKAAGKYRQQGRNY
IVEDGDI IFFKFNT PQPKKK

971/6881
FIGURE 905

CTGCCAGCCGCGCTGCTGCTGCTCCTCCTGCTGTGGGACCGCTGACCGCGCGGCTGCTCCGCTCTCCCCGCTCCA
AGCGCCGATCTGGGCACCCGCCACCAGCATGGACGCTCGCCGCGTGCCGCAGAAAGATCTCAGAGTAAAGAAGAA
CTTAAAGAAATTCAGATATGTGAAGTTGATTTCCATGGAGACCTCGTCATCCTCTGATGACAGTTGTGACAGCTT
TGCTTCTGATAATTTTGCAAACACGAAACCTAAATTCAGGTGAGATATCAGTGAAGAACTGGCAAGTGTTTTTTA
TGAGGACTCTGATAATGAATCTTTCTGCGGCTTTTCAGAAAGTGAGGTGCAAGATGTATTAGACCATTGTGGATT
TTTACAGAAACCAAGGCCAGATGTCTACTAACGAACCTGGCCGGTATTTTTCATGCCGACTCTGACGATGAATCATT
TTGCGGTTTCTCAGAGAGTGAGATACAAGATGGAATGAGGCTGCAGTCAGTTCGGGAAGGCTGTAGGACCCGCAG
CCAGTGACAGGCACTCTGGACCTCTCAGGGTGGCGATGAAGTTTCCAGCGCGGAGTACCAGGGGAGCAACCAACAA
AAAAGCAGAGTCCC GCCAGCCCTCAGAGAATTCTGTGACTGATTCCAACCTCCGATTGAGAAGATGAAAGTGGAAT
GAATTTTTTGGAGAAAAGGGCTTTAAATATAAAGCAAACAAAGCAATGCTTGCAAAACTCATGTCTGAATTAGA
AAGCTTCCCTGGCTCGTTCCGTGGAAGACATCCCCTCCCAGGCTCCGACTCACAATCAAGGAGACCGCGAAGGCG
TACATTTCCCGGTGTTGCTTCCAGGAGAAACCTGAACGGAGAGCTCGTCTCTTACCAGGTCAGGTCCCAGAT
CCTCGGGTCCCCTTGACGCTCTACCCATGGAGGAGGAGGAGGAAGAGGATAAGTACATGTTGGTGAGAAAGAGGAA
GACCGTGGATGGCTACATGAATGAAGATGACCTGCCCAGAAGCCGTCGCTCCAGATCATCCGTGACCCCTTCCGCA
TATAATTCGCCCAGTGGAAGAAATTACAGAGGAGGAGTTGGAGAACGCTGTCAGCAATTCTCAGAGAGAAGATATA
TAACCGTTCACTGGGCTCTACTTGTCTCATCAATGCCGTCAGAAGACTATTGATACCAAAACAACTGCAGAAACCC
AGACTGCTGGGGCGTTTCGAGGCCAGTTCTGTGGCCCTGCCTTCGAAACCGTTATGGTGAAGAGGTCAGGGATGC
TCTGCTGGATCCGAACCTGGCATTGCCCGCCTGTGCGAGGAATCTGCAACTGCAGTTTCTGCCGCCAGCGAGATGG
ACGGTGTGCGACTGGGGTCTTGTGTATTTAGCCAAATATCATGGCTTTGGGAATGTGCATGCCCTACTTGAATAAG
CCTGAAACAGGAATTTGAAATGCAAGCATAATATCTGGAATAATTTGCTGCCTGCCTTCTACTTCTCAAATCTTTC
TTGTAAAGTTTCCAATTTTTTCACTGAAACCTGAGTTAAAAATCTTGATGATCAGCCTGTTTCAATAAGAACTC
CAATCAAGTTAATCTTAGCAGACATGTGTTTCTGGAGCATCAGAGAAGGTATATTGCTAGTTACACTTTGCCCTC
CTGCAGTTTCTTCTGCTCCCAACCCCATCTCATAGCATCCCCCTCTATTTCCAATGCTCCTCTCCAACCGCT
TAGTTTCTGAATTTCTTTAAATTACAGTTTTATGAAAGCATATTTTATTTACTTGGTGTTGAAATAGCCCTCAT
AAAACCTAAGCACTTGGAAACACAATAATAGTATTAACCTAAGTATCTATTGAATTTGAGAGAAGACCTTCTA
ACTTGTTTACACAAAAACGAGTATGATTTAGCATTCTACTAGTTGAAATTTTTAATAGAATCAAGGCACAAAAG
TCTTAAACCATGTGGAAAAATTAGGTAATTATTCAGATTGATGTCCTCAATCCCATGTATTGCGCTTATGTT
ACAAGTTGTTGTACAGTTGAGACTTAATTTCTCCTAATTTCTTCTGCCCGAAGGGTAAGTGGTGCGTCCAGCTT
ACACAATCATAATTCAAAGGTTGGTGGGCAATGTAATACCTAATTAATAAATAATGATGGAAGAGCTATCTGGAGAT
TATGAGTAAGCTGATTTGAATTTTTCAGTATAAACTTTAGTATAATTGTAGTTTGCAAAGTTTATTTAGTTTAC
ATGTAAGGTATTGCAAATAAATCTTGGACAATTTTGTATGGAACCTTGATATTAAAACTAGTCTGTGGTTCTT
TGCAGTTTCTTGTAATTTATAAACCAGGCACAAGGTTCAAGTTTAGATTTTAAGCACTTTTATAACAATGATAA
GTGCCTTTTTGGAGATGTAACTTTTAGCAGTTTGTAACTGACATCTCTGCCAGTCTAGTTTCTGGGCAGGTTT
CCGTGTGTCAGTATTTCCCTCCTCTTTGCATTAAATCAAGGTATTTGGTAGAGGTGGAATCTAAGTGTGTTGTATGT
CCAATTTACTTGATATGTAACCATTTGCTGTGCTATTCAATGTTTGATGCATAATTGGACCTTGAATCGATAAG
TGTAATAACAGCTTTTGATCTGTAATGCTTTTATACAAAAGTTTATTTTAATAATAAAATGTTTGTCT

972/6881
FIGURE 906

MDARRVPQKDLRVKKNLKKFRYVKLISMETSSSSDDSCDSFASDNFANTKPKFRSDISEELASVFYEDSDNESFC
GFSESEVQDVLHDHCGFLQKPRPDVTNELAGIFHADSDDESFCGFSESEIQDGMRLQSVREGCRTRSQCRHSGPLR
VAMKFFPARSTRGATNKKAESRQPSSENSVTDSNSDSEDESGMNFLEKRALNIKQNKAMLAKLMSELESFPGSFRGR
HPLPGSDSQSRRPRRRTFFPGVASRRNPERRARPLTRSRILGSLDALPMEEEEEEDKYMLVRKRKTVDGYMNED
DLPRSRSSSVTLPHIIRPVEEITEELENVCSNSREKIYNRSLGSTCHQCRQKTIDTKTNCRNPD CWGVRGQF
CGPCLRNRYGEEVRDALLDPNWHCPPCRGICNCSCFCRQDGRCATGVLVYLAKYHGFGNVHAYLKSLKQEFEMQA

973/6881
FIGURE 907

GCAAGATGGCAGAAGTAGAGCAGAAGAAGAAGCGGACCTTCGCAAGTTCACCTACCGCGGCGTGGACCTGGACC
AGCTGCTGGACATGTCCTACGAGCAGCTGGAGCAGCCGATGCAGCTGTACAGTGC GCGCCAGCGGCGGCGGCTGA
ACCGGGGCCTGCGGCGGAAGCAGCACTCCCTGCTGAAGCGCCTGCGCAAGGCCAAGAAGGAGGCGCCGCCCATGG
AGAAGCAGGAAGTGGTGAAGACGCACCTGCGGGACATGATCATCCTACCCGAGATGGTGGGCAGCATGGTGGGCG
TCTACAACGGCAAGACCTTCACCTACAAGCCCGTAAAGCACGGCCGGCCCGGCATCGGGGCCACCCACTCCTCCC
GCTTCATCCCTCTCAAGTAAATGGCTCAGCTAATAAAGGCGCACATGACTCC

974/6881

FIGURE 908

MAEVEQKKKRTFRKFTYRGVDLDQLDMSYEQLEQPMQLYSARQRRRLNRGLRRKQHSLLKRLRKAKKEAPPMEK
QEVVKTHLRDMIILPEMVGSMVGVYNGKTFTYKPVKHGRPGIGATHSSRFIPLK

975/6881
FIGURE 909

GCTGAAATTAAAAGTACTGAGCTTACCAAGTGTGATAAGGATGTAGAGGAAC TGGAAC TTT CATACACTGCTGG
TGGGAATATAAAATGGTATAACCATTTTGGAATACAAC TTGGCAGTTTATTTGTTTGT TTAGGTTGTTTTACTC
ACCACTATGCTAACGAGGAACTTGGCAGTTTCTTTAAAAGATAAACATATATCTCTGGTCCGTGCCTCCAAGATG
ACAAAGAAAAGAAGGAACAACGGTCATGCCAAAAAGGGCCGCGGCCACGTGCATCCTATT CGCTGCACTAACTGT
ACCCGATGCGTGCCCAAGGACAAGGCCATTAAGAAATTCGTCATT CGAAACGTAGTGGAGGCCG CAGCAGTCAGG
GACATTCTGAAGTGAGCGTCITCAATGCCTATGTGCTTCCCAAGCTGTATGTGAAGCTATATTACTGTGTGAGT
TGTGCAATTCACAGCAAAGTAGTCAGGAATCGATCTCGTGAAGCC CACAAAGACCGAACACCCCCACCTGATTT
AGACCTGTGGGTGCTGCCCCACGTCCCCACCAAAGCCCATGTAAGGAGCTGAGTTCTTAAAGACTGAAGACAGA
CTATTCTCTGGAGAAAAATAAAATGGAAATTGTACTT

976/6881
FIGURE 910

CTTTTATGTTAGATGAGGAAGGGGATACCCAAACAGAGGAAACCCGGCCTTCAGAAACAAAAGAAGTGGAGCCAG
AGCCAACTGAGGACAAAGATTTGGAAGCTGATGAAGAGGACACTAGGAAAAAAGATGCTTCTGATGATCTAGATG
ACTTGAACCTCTTTAATCAAAAGAAAAAGAAGAAAAAACTAAAAAGATATTTGATATTGATGAAGCTGAAGAAG
GTGTAAAGGATCTTAAGATTGAAAGTGATGTTCAAGAACCAACTGAACCAGAGGATGACCTTGACATTATGCTTG
GCAATAAAAAGAAGAAAAAGAAGAAATGTTAAGTTCCCAGATGAGGATGAAGCTCTAGAAGATGAAGACAACAAA
AAGATGATGGTATCTCATTCAAGTAATCAGACAGGCCCTGCTTGGGCAGGCTCAGAAAGAGACTACACATACGAGG
AGCTGCTGAATCGAGTGTCAACATCATGAGGGAAAAGAATCCAGATATGGTTGCTGGGGAGAAAAGGAAATTTG
TCATGAAACCTCCACAAGTCGTCCGAGTAGGAACCAAGAAAACCTTCTTTTGTCAACTTTACAGATATCTGTAAAC
TTGGTTCTATAGATGGTAATAACCAACTTGTAATCAAAGGAAGATTCCAACAGAAACAGATAGAAAATGTCTTGA
GAAGATATATCAAGGAGTATGTCACTTGTACACATGCCGATCACCGACACAATCCTGCAGAAGGACATGCGAC
TCTATTTCTACAGTGCAGAACTTGTCACTTAGATGTTCTGTTGCCATTATCAAATTTGGCTTCCAGGCTGTCA
TGGGCAAGCGAGCACAGCTCCGTGCCAAAGCTAACTAATTTGCTAATCACTGATTTTGCAAAGCTTGTGTGGAG
ATGTGGCTGGACAGGTTTGCCATCAGAGTGGATATAACATTGTATTAAAAACAAGATAAAAAAGCTGCCAAGATT
TTTGGCGAGTGGTTGGTTGGTCTGAAGTCCTTGCAAGACGCTGATGCTCAAGCTGTTGACATACTCATGCTTAC
TTTAACACCTGTCAGAGAAACGTGATATGGGGTAAGGAGGTGCTTTTTTAAATAGTTCATAGACTTCTGTAAAA
TGCAAGATAAATTAAAGTTATTATAACAGTGA

977/6881
FIGURE 911

GGTGGGAGAGGTATCGGCAGGGGCAGCGCTGCCGCCGGGGCCTGGGGCTGACCCGTCTGACTTCCCGTCCGTGCC
GAGCCCACTCGAGCCGCAGCCATGTCTGGGGACGAGATGATTTTTGATCCTACTATGAGCAAGAAGAAAAAGAAG
AAGAAGAAGCCTTTTATGTTAGATGAGGAAGGGGATACCCAAACAGAGGAAACCCGGCCTTCAGAAACAAAAGAA
GTGGAGCCAGAGCCAACTGAGGACAAAGATTTGGAAGCTGATGAAGAGGACACTAGGAAAAAAGATGCTTCTGAT
GATCTAGATGACTTGAACCTCTTTAATCAAAAGAAAAAGAAGAAAAAACTAAAAAGATATTTGATATTGATGAA
GCTGAAGAAGGTGTAAAGGATCTTAAGATTGAAAGTGATGTTCAAGAACCAACTGAACCAGAGGATGACCTTGAC
ATTATGCTTGGCAATAAAAAGAAGAAAAAGAAGATGTTAAGTTCCCAGATGAGGATGAAGCTCTAGAAGATGAA
GACAACAAAAAGATGATGGTATCTCATTCAAGTAATCAGACAGGCCCTGCTTGGGCAGGCTCAGAAAGAGACTAC
ACATACGAGGAGCTGCTGAATCGAGTGTTCAACATCATGAGGGAAAAGAATCCAGATATGGTTGCTGGGGAGAAA
AGGAAATTTGTCATGAAACCTCCACAAGTCGTCCGAGTAGGAACCAAGAAAACCTCTTTTGTCAACTTTACAGAT
ATCTGTAAACTATTACATCGTCAACCCAAACATCTCCTTGCAATTTTATTGGCTGAATTGGGTACAAGTGTTCT
ATAGATGGTAATAACCAACTTGTAATCAAAGGAAGATTCACACAGAAACAGATAGAAAATGTCTTGAGAAGATAT
ATCAAGGAGTATGTCATTGTACACATGCCGATCACCGGACACAATCCTGCAGAAGGACATGCGACTCTATTTT
CTACAGTGCAGAACTTGTCAATCTAGATGTTCTGTTGCCATTATCAAAATTGGCTTCCAGGCTGTCATGGGCAAG
CGAGCACAGCTCCGTGCCAAAGCTAACTAATTTGCTAATCACTGATTTTGCAAAGCTTGTTGTGGAGATGTGGCT
GGACAGGTTTGCCATCAGAGTGGATATACCATGTATTAAAAACAAGATAAAAAAGCTGCCAAGATTTTGGCGA
GTGGTTGGTTGGTCTGAAGTCCTTGCAAGACGCTGATGCTCAAGCTGTTGACATACTCATTGCCTACTTTAACAC
CTGTCAGAGAAACGTGATATGGGGTAAGGAGGTGCTTTTTTAAATAGTTTCATAGACTTCTGTAAATGCAAGAT
AAATTAAAGTTATTATAACAGTGA

978/6881
FIGURE 912

CGAGTTGGAAGAGGCGAGTCCGGTCTCAAAATGGAGGTAAAACCGCCGCCCGGTGCGCCCCAGCCCGACTCCGGC
CGTCGCCGTGCGCCGCCGGGGGGAGGAGGGCCATGATCCAAAGGAACCAGAGCAGTTGAGAAAACGTGTTTATTGGT
GGTCTGAGCTTTGAAACTACAGATGATAGTTTACGAGAACATTTTGAGAAATGGGGCACACTCACAGATTGTGTG
GTAATGAGAGACCCCCAAACAAAACGTTCCAGGGGCTTTGGTTTTGTGACTTATTCTTGTGTTGAAGAGGTGGAT
GCAGCAATGTGTGCTCGACCACACAAGGTTGATGGGCGTGTAGTGGAACCAAAGAGAGCTGTTTCTAGAGAGGAT
TCTGTAAAGCCTGGTGGCCATCTAACAGTGAAGAAAATTTTGTGTTGGTGGTATTAAAGAAGATACAGAAGAATAT
AATTTGAGAGACTACTTTGAAAAGTATGGCAAGATTGAAACCATAGAAGTTATGGAAGACAGGCAGAGTGGAAAA
AAGAGAGGATTTGCTTTTGTAACTTTGTATGATCATGATACAGTTGATAAAATTGTTGTTTCAGAAAATACCACACT
ATTAATGGGCATAATTGTGAAGTGAAAAAGGCCCTTTCTAAACAAGAGATGCAGTCTGCTGGATCACAGAGAGGT
CGTGAGGTTGGATCTGGCAATTTTATGGGTGCGCGGAGGGAACTTTGGAGGTGGTGGAGGTAATTTTGGCCGTGGT
GGAAACTTTGGTGGAGAGGTAGGCTATGGTGGTGGAGGTGGTGGCAGCAGAGGTAGTTATGGAGGAGGTGATGG
TGATATAATGGATTGGAGGTGATGGTGGCAACTATGGCGGTGGTCTGTTATAGTAGTAGAGGGGGCTATGG
TGGTGGTGGACCAGGATATGGAACCAAGGTGGTGGATATGGTGGAGGTGGAGGATATGATGGTTACAATGAAGG
AGGAAATTTTGGCGGTGGTAACTATGGTGGTGGTGGGAACATAATGATTTTGGAAATTATAGTGGACAACAGCA
ATCAAATTATGGACCCATGAAAGGGGGCAGTTTGGTGGAGAAGCTCGGGCAGTCCCTATGGTGGTGGTTATGG
ATCTGGTGGTGAAGTGGTGGATATGGTAGCAGAAGGTTCTAAAAACAGCAGAAAAGGGCTACAGTTCTTAGCAG
GAGAGAGAGCGAGGAGTTGTCAGGAAAGCTGCAGGTTACTTTGAGACAGTCGTCCCAAATGCATTAGAGGAACTG
TAAAAATCTGCCACAGAAGGAACGATGATCCATAGTCAGAAAAGTTACTGCAGCTTAAACAGGAAACCCCTTCTTG
TTCAGGACTGTCATAGCCACAGTTTGCAAAAAGTGCAGCTATTGATTAATGCAATGTAGTGTCAATTAGATGTAC
ATTCTGAGGTCTTTTATCTGTTGTAGCTTTGTCTTTTTCTTTTTCTTTTTCATTACATCAGGTATATTGCCCTGT
AAATTGTGGTAGTGGTACCAGGAATAAAAAATTAAGGAATTTTAACTTTTCAATATTTGTGTAGTTCAGTTTTT
CTACATTTTAGTACAGAACTTTAACAAAATGCAGTTTCGAAGGTGTTTCCTTGTGAGTTAACAAGTAAAGAAGA
TCATTGTTAATTACTATTTTGTATGAATTTTGCTAAAGTTAACTGTAAAGAAACACCTGCTGACTTGCAGTTTAA
GGGAATCTATTCTCCCATTTCCAAACCATGATATGAATGGGCGCTGACATGTGGAGAGAATAGATAATTTGTG
TGTTTGCAATGTGTGTTTGTAGATAAATAGGATTGGGTATTTAAATTAGCATTGTGAATTTAATAGCATTAAAGAT
TACCTTCAAATGAAAAAAATCTCAAAATTTCTATTTGGTTTTGTGCATTTCTTTTAAATGTAATCATATGA
TTTTAGTGTGTTAGACTTGCTGAGTCCTAGCTGTGTTTAGAACATCTCTATTCTACATTTACCTTGGTCAAATTT
GAACTGCTGCCATAGG

979/6881
FIGURE 913

MEGHDPKEREQLRKFFIGGLSFETITDDGLREHFKEKWWTLTDCVVMRDPQTKCSRGFGFVTYSYIEEVDAAMCAPP
HKVDGCVVEPKRAVSREDFVKPGAHLTVKKIFVGGIKIQKNII

980/6881
FIGURE 914

CTTCATCCTGCGACTAGCACCGCGTCCGGCAGCGCCTGCCCTACACTCGCCCGTGCCATGGCCTCCGTCTCCAAG
CTCGCCTGCATCTACTCGGCCCTCATTCTGCACGACGATGAGGTGACAGTCACGGAGGATAAGATCAATGCCCTC
ATTAAAGCAGCCGGTGTAATGTTGAACCTTTTTGGCCTGGCTTGTTTGCAAAGGCCCTGGCCAACGTCAACATT
GGGAGCCTCATCTGCAATGTAGGGGCTGGTGGACCTGCTCCAGCTGAGGAGAAGAAAGTGGGAAGCAAAGAAAGAA
GAATCCAAGGAGTCTGATGATGACATGGGCCTTGGTCTTTTGGACTAAACCTCTTTTATAACGTGTTCAATAAAA
AGCTGAACCT

981/6881
FIGURE 915

MASVSKLACIYSALILHDDEVTVTEDKINALIKAAGVNVEPFWPGLFAKALANVNIGSLICNVGAGGPAPAEKK
VEAKKEESKESDDDMGLGLFD

982/6881
FIGURE 916

ACGGGGCCTGGGCGGSAGGGGCGGTGGCTGGAGCTCGGTAAAGCTCGTGGGACCCCATTTGGGGGAATTTGATCCA
AGGAAGCGGTGATTGCCGGGGGAGGAGAAGCTCCCAGATCCTTGTGTCCACTTGCAGCGGGGGAGGCGGAGACGC
GGAGCGGGCCTTTTGGCGTCCACTGCGCGGCTGCACCCTGCCCCATCCTGCCGGGATCATGCTCTGCGGCAGCCC
GGGAGGGATGCTGCTGCTGCGGGCCGGGCTGCTTGCCCTGGCTGCTCTGCTGCTCCGGGTGCCCGGGGCTCG
GGCTGCAGCCTGTGAGCCCGTCCGCATCCCCCTGTGCAAGTCCCTGCCCTGGAACATGACTAAGATGCCCCAACCA
CCTGCACCACAGCACTCAGGCCAACGCCATCCTGGCCATCGAGCAGTTCGAAGGTCTGCTGGGCACCCACTGCAG
CCCCGATCTGCTCTTCTTCTCTGTGCCATGTACGCGCCCATCTGCACCATTGACTTCCAGCACGAGCCCATCAA
CCCTGTAAAGTCTGTGTGCGAGCGGGCCCGGCAGGGCTGTGAGCCCATACTCATCAAGTACCGCCACTCGTGGCC
GGAGAACCTGGCCTGCGAGGAGCTGCCAGTGTACGACAGGGGCGTGTGCATCTCTCCCGAGGCCATCGTTACTGC
GGACGGAGCTGATTTTCCTATGGATTCTAGTAACGGAACTGTAGAGGGGCAAGCAGTGAACGCTGTAAATGTAA
GCCTATTAGAGCTACACAGAAGACCTATTTCCGGAACAATTACAACATATGTCATTGGGGCTAAAGTTAAAGAGAT
AAAGACTAAGTGCCATGATGTGACTGCAGTAGTGGAGGTGAAGGAGATTCTAAAGTCTCTCTGGTAAACATTCC
ACGGGACACTGTCAACCTCTATACCAGCTCTGGCTGCCTCTGCCCTCCACTTAATGTTAATGAGGAATATATCAT
CATGGGCTATGAAGATGAGGAACGTTCCAGATTACTCTTGGTGAAGGCTCTATAGCTGAGAAGTGAAGGATCG
ACTCGGTAAAAAAGTTAAGCGCTGGGATATGAAGCTTCGTCATCTTGGACTCAGTAAAAGTGATTCTAGCAATAG
TGATTCCACTCAGAGTCAGAAGTCTGGCAGGAACTCGAACCCCGGCAAGCACGCAACTTAAATCCCGAAATACAA
AAAGTAACACAGTGGACTTCCTATTAAGACTTACTTGCATTGCTGGACTAGCAAAGGAAAATTGCACTATTGCAC
ATCATATTCTATTGTTTACTATAAAAATCATGTGATAACTGATTATTACTTCTGTTTCTCTTTTGGTTTCTGCTT
CTCTCTTCTCTCAACCCCTTTGTAATGGTTTGGGGGCAGACTCTTAAGTATATTGTGAGTTTTCTATTTCACTAA
TCATGAGAAAACTGTTCTTTTGCAATAATAATAAATTAAACATGCTGTTA

983/6881

FIGURE 917

MVCGSPGGMLLLRAGLLALAALCLLRVPGARAAACEPVRIPLCKSLPWNMTKMPNHLHHSTQANAILAIEQFEG
LGTHCSPDLLFFLCAMYAPICTIDFQHEPINPCKSVCEARQGCEPILIKYRHSWPENLACEELPVYDRGVCISP
EAIVTADGADFPMDSNGNCRGASSERCKCKPIRATQKTYFRNNYNYVIRAKVKEIKTKCHDVTAVVEVKEILKS
SLVNIPRDTVNLYTSSGCLCPPLNVNEEYIIMGYEDEERSRLLLVEGSIAEKWKDRLGKKVWRWDMKLRHLGLSK
SDSSNSDSTQSQKSGRNSNPRQARN

984/6881

FIGURE 918

ATGTCCATCAGGACGACCCAGAAGTCCTACAGGGTGTCCACTTCTGGCCCCCGGGCCTTCAGCAGCCATTTCTAC
ACAAGTGGGCCTGGTGCCTCCATCAGCTCCTCGAGCTTCTCCCAAGTGGGCAGCAGCAGCTTCCGGGGTGGCCTG
GGAGGAGGCTACGGTGGGGCCAGTGGCATGGGAGGCATCACCACCGTCACTGTCAACCAGAGCCTGCTGAGCCCC
CTTAACCTGGAGGTGGACCCACACATCCAGGCAGTGCACTCAGAAGAAGGAGCAGATCAAGACCCTCAACACT
AAGTTTGCCACCTTCATATACAAGAGTTACATCAACAACCTTAGGCAGCAGCTGGAGACTCTGGGCCAGGAAAAG
CTGAAGCTGGAGGCGGAGCTTGGAACATGCAGGGGCTGGTGGAGGACTTCAAGAACAAGTATGAGGATGAGATC
AATAAGCGTACAGAGATGGAGAATGAATTTGTCTCATCAAGAAGGATGTGGATGAAGCTTACCTGAACAAGGTA
GAGCTGGAGTCTCGCCTGGAAGGGCTGACTGACGAAATCAACTTCCTCAGGCAGCTGTATGAAGAGGATATCCCG
GAGCTGCAGTCCCAGATCTCAGACACGTCTGTGGTGCTGCCCATGGACAACAGCCACTCCCTGGACATGGACAGC
ATCATCAATGAGGTCAAGCGCAGTACGAGATCACCAACCGCAGCCAGCTGGCTCCAGGCAGAGATTGAGGGTCTC
AAAGGCCAGAGGGCTTCCCTGGAGGCCGCCATCGCAGATGCGGAGCAGCGCGGGAGTTGGCCGTTAAGGATGCC
AGCGCCAAGCTGTCTGAGCTGGAGGCCGCCCTGCAGCAAACCAAGCAGGACGTGGAGCTGATGAACGACAAGCTG
GCCCTGGACATTGAGATCGCCACCTACAGGCAGCTGCTGGAGGGCGAGGAGAGCTGGCTGGAGTCTGGGATGCAG
AGCATGAGTATCCATACAAAGACCATCAGCAGCTATGCAGGTGGTCTGAGCTCGGCCTATGGGGGCCTCACAAAGC
CCCGCCTCAGCTGTGGACTGGGCTCCAGCTTTGGCTCTGGCGGGGCTCCAGTTCCTTCAGCTGCATCAGCTAC
ACCAGGGCCGTGGTTGTGAAGAAGATTGAGACCCGTGATGGGACGCTGGTGTGAGAGTTCTCTGACGTCCTGCCC
AAGTGA

985/6881
FIGURE 919

MSIRTTQKSYRVSTSGPRAFSSHFYTSGPGASISSSSFQVGSSSFRRGGLGGGYGGASGMGGITTVTVNQSLSP
LNLEVDPHIQAVHTQKKEQIKTLNFKFATFIYKSYINNLRQQLTLGQEKLEAELGNMQGLVEDFKNKYEDEI
NKRTEMENEFVLIKKDVDEAYLNKVELESRLGLTDEINFLRQLYEEDIPELQSQISDTSVVLPMDNSHSLDMS
IINEVKRSTRSPTAASWLQAEIEGLKGQRASLEAAIADAEQORGELAVKDASAKLSELEAALQQTQDVELMNDKL
ALDIEIATYRQLLEGEESWLESGMQSMSIHTKTISSYAGGLSSAYGGLTSPGLSCGLGSSFGSGAGSSSFSCISY
TRAVVVKKIETRDGTLVSEFSDVLPK

986/6881

FIGURE 920

ACGCAACCCCATAGGTAGCACGCCGTTACCCGTGGGGAGCGTTTCCGCCATTTTGGAAAATTAATTGGGAAGGT
ACTGGTTTTAAGTGTAGTTGCCGACGCAATGGCAGCCTTTGCAGTGGAACCTCAGGGGCGCGTTAGGATCTGA
ACCAATGATGCTGGGTTACCCACATCTCCAAAGCCAGGAGTTAATGCCAGTTCTTACCTGGATTTTAAATGGG
GGATTTGCCAGCTCCGGTGACTCCACAACCTCGATCAATTAGTGGCCCTTCAGTAGGAGTAATGGAAATGAGATC
ACCTTTACTTGCAGGTGGGTCAACCACCACAACCAGTTGTACCAGCTCATAAAGATAAAAGTGGCGCTCCACCAGT
TAGAAGTATATATGATGACATTCTAGCCCAGGACTTGGATCAACACCTTTAACTTCAAGAAGACAGCCAAACAT
TTCAGTAATGCAGAGTCTCTTGTGGAGTTACATCTACTCCTGGAACAGGGCAAAGTATGTTTAGTCCAGCAAG
TATCGGTGAGCCAGCAAGACGACATTATCTCCTGCCAGTTGGATCCTTTTTATACTCAAGGAGATTCTTTGAC
TTCAGAAGATCACCTCGATGACTCTTGGGTGACTGTATTTGGGTTTCTCAAGCATCTGCTTCTACATATTACT
ACAATTTGCACAGTATGGGAATATCTTAAACATGTGATGTCTAATACAGGAAATTGGATGCATATTCGTTATCA
ATCTAAACTGCAGGCTCGGAAAGCCTTAAGCAAAGATGGGAGGATTTTGGAGAATCCATCATGATTGGTGTA
ACCATGTATTGACAAAAGTGTTATGGAAAGCAGTGACAGATGTGCTTTATCATCTCCATCTTTAGCCTTTACACC
ACCAATCAAACTCTAGGTACACCAACAACCTGGAAGTACTCCTAGGATTTCTACCATGAGACCTCTTGCTAC
AGCATACAAAGCCTCTACTAGTGATTATCAGGTTATTTCTGACAGACAAACGCCAAAAAAGATGAAAGTCTTGT
ATCCAAAGCAATGGAGTACATGTTTGGCTGGTAGTAGAACACCAAGAAGGAGGTGCTACACTAAAACAGAGTTA
GCAGAGTGCTGCTGGTTTCCTTCGGTTAGTTATATACTGTTCTGCGAGTATTGGATAGCTATCTCATACTTCTTT
TAGAAAGAAGCCTTTTTTATTAAAGGATACAACCTATTTGTAGCTCGCACTTTAAAGATGCTTGAGATACATTTT
AAAGAAAACATAAAATCCCTGTAAATAGGATTTTGTGCTTTCTGTAAACAGTGCATGCTTCAGCACAGAAAAC
GCATTGATTATTGTAAATTAAATACTGAAATTGTGGTGAGACGTCATAGTCTTCATGAGAACGTGGGGGTGAAT
TTCATGAAGGGGAACATATAGTTATTTCTACCGACACAAATATTATAATTAGCAATTTGAATTATGGTCTTTTAA
TTAGATAGTATTTAATATTTAATTATCCTTGTTTGTATATGTCCTGTCACAGAGTGTCCTCTTGGTGTATTCTA
AAACGAGCATTTCTTTAAAAAACCTAAAGTTTCTTGATAATAAACATTGTCAATGAT

987/6881
FIGURE 921A

GGCTGAGTTTTATGACGGGCCCCGGTGCTGAAGGGCAGGGAACAACCTTGATGGTGCTACTTTGAACTGCTTTTCTT
TTCTCCTTTTTTGCACAAAGAGTCTCATGTCTGATATTTAGACATGATGAGCTTTGTGCAAAAGGGGAGCTGGCTA
CTTCTCGCTCTGCTTCATCCCACTATTATTTTGGCACAAACAGGAAGCTGTTGAAGGAGGATGTTCCCATCTTGGT
CAGTCTATGCGGATAGAGATGTCTGGAAGCCAGAACCATGCCAAATATGTGTCTGTGACTCAGGATCCGTTCTC
TGCGATGACATAATATGTGACGATCAAGAATTAGACTGCCCAACCCAGAAATTCATTGGAGAATGTTGTGCA
GTTTGGCCACAGCCTCCAACTGCTCCTACTCGCCCTCCTAATGGTCAAGGACCTCAAGGCCCAAGGGAGATCCA
GGCCCTCCTGGTATTCTGGGAGAAATGGTGACCCTGGTATTCCAGGACAACCAGGGTCCCCCTGGTTCTCCTGGC
CCCCCTGGAATCTGTGAATCATGCCCTACTGGTCTCAGAACTATTCTCCCCAGTATGATTATATGATGTCAAG
TCTGGAGTAGCAGTAGGAGGACTCGCAGGCTATCCTGGACCAGCTGGCCCCCAGGCCCTCCCGGTCCCCCTGGT
ACATCTGGTCATCCTGGTTCCCTGGATCTCCAGGATACCAAGGACCCCCCTGGTGAACCTGGGCAAGCTGGTCCT
TCAGGCCCTCCAGGACCTCCTGGTGCTATAGGTCCATCTGGTCTGCTGGAAAAGATGGAGAATCAGGTAGACCC
GGACGACCTGGAGAGCGAGGATTGCCTGGACCTCCAGGTATCAAAGGTCCAGCTGGGATACCTGGATTCCCTGGT
ATGAAAGGACACAGAGGCTTCGATGGACGAAATGGAGAAAAGGGTGAAACAGGTGCTCCTGGATTAAAGGGTGAA
AATGGTCTTCCAGGCGAAAATGGAGCTCCTGGACCCATGGGTCCAAGAGGGGCTCCTGGTGAGCGAGGACGGCCA
GGACTTCTGGGGCTGCAGGTGCTCGGGGTAATGACGGTGCTCGAGGCAGTGATGGTCAACCAGGCCCTCCTGGT
CCTCCTGGAATGCGCGATTCCCTGGATCCCTGGTGCTAAGGGTGAAAGTTGGACCTGCAGGGTCTCCTGGTTCA
AATGGTGCCCTGGACAAAGAGGAGAACCTGGACCTCAGGGACACGCTGGTGCTCAAGGTCTCCTGGCCCTCCT
GGGATTATGGTAGTCTTGGTGGTAAAGGCGAAATGGGTCCCGCTGGCATTCTGGAGCTCCTGGACTGATGGGA
GCCCCGGGTCTCAGGACCAGCCGGTGCTAATGGTGCTCCTGGACTGCGAGGTGGTGAGGTGAGCCTGGTAAG
AATGGTGCCAAAGGAGAGCCCCGACCACGTGGTGAACGCGGTGAGGCTGGTATTCCAGGTGTTCCAGGAGCTAAA
GGCGAAGATGGCAAGGATGGATCACCTGGAGAACCTGGTGCAAATGGGCTTCCAGGAGCTGCAGGAGAAAGGGT
GCCCCTGGGTTCAGGAGCTGCTGGACCAAATGGCATCCAGGAGAAAAGGGTCTGCTGGAGAGCGTGGTGCT
CCAGGCCCTGCAGGGCCCAGAGGAGCTGCTGGAGAACCTGGCAGAGATGGCGTCCCTGGAGGTCCAGGAATGAGG
GGCATGCCCGGAAGTCCAGGAGGACCAGGAAGTGATGGGAAACCAGGGCCTCCCGGAAGTCAAGGAGAAAGTGGT
CGACCAGGTCTCCTGGGCCATCTGGTCCCCGAGGTGAGCTGGTGCTCATGGGCTTCCCCGGTCTTAAAGGAAAT
GATGGTGCTCCTGGTAAGAAATGGAGAACGAGGTGGCCCTGGAGGACCTGGCCCTCAGGGTCTCCTGGAAAGAAAT
GGTGAACTGGACCTCAAGGACCCCCAGGGCTACTGGGCTGGTGGTGACAAAGGAGACAGGACCCCTGGT
CCACAAGGATTACAAGGCTTGCTGGTACAGGTGGTCTCCAGGAGAAAATGGAAAACCTGGGGAACCAAGGTTCA
AAGGGTGATGCCGGTGACCTGGAGCTCCAGGAGGCAAGGGTGATGCTGGTGCCCTGGTGAACGTGGACCTCCT
GGATTGGCAGGGGCCCCAGGACTTAGAGGTGGAGCTGGTCCCCCTGGTCCCGAAGGAGGAAAGGGTGCTGCTGGT
CCTCCTGGGCCACCTGGTGCTGCTGGTACTCCTGGTCTGCAAGGAATGCCTGGAGAAAGAGGAGGTCTTGGAAAT
CCTGGTCCAAAGGGTGACAAGGGTGAAACAGGCGGCCAGGTGCTGATGGTGCTCCAGGGAAGATGGCCCAAGG
GGTCTACTGGTCTATTGGTCTCCTGGCCAGCTGGCCAGCCTGGAGATAAGGGTGAAGGTGGTGGCCCCGGA
CTTCCAGGTATAGCTGGACCTCGTGGTAGCCCTGGTGAGAGAGGTGAAACTGGCCCTCCAGGACCTGCTGGTTTC
CCTGGTGCTCCTGGACAGAATGGTGAACCTGGTGGTAAAGGAGAAAGAGGGGCTCCGGGTGAGAAAGGTGAAGGA
GGCCCTCCTGGAGTTGACAGGACCCCTGGAGGTTCTGGACCTGCTGGTCTCCTGGTCCCCAAGGTGTCAAAGGT
GAACGTGGCAGTCTTGGTGGACCTGGTGCTGCTGGCTTCCCTGGTGCTCGTGGTCTTCTGGTCTCCTGGTAGT
AATGGTAACCCAGGACCCCAAGTCCCAGCGGTTCTCCAGGCAAGGATGGGCCCCCAGGTCTGCGGGTAACACT
GGTGCTCCTGGCAGCCCTGGAGTGCTGGACCAAAGGTGATGCTGGCCAACCAGGAGAGAAGGGATCGCCTGGT
GCCCAGGGCCCACAGGAGCTCAGGCCCACTTGGGATTGCTGGGATCACTGGAGCACGGGGTCTTGCAGGACCA
CCAGGCATGCCAGGTCTAGGGGAAGCCCTGGCCCTCAGGGTGTCAAGGGTGAAAGTGGGAAACCAGGAGCTAAC
GGTCTCAGTGGAGAACGTGGTCCCCCTGGACCCAGGGTCTTCTGGTCTGGCTGGTACAGCTGGTGAACCTGGA
AGAGATGGAAACCTGGATCAGATGGTCTTCCAGGCCGAGATGGATCTCCTGGTGGCAAGGGTGATCGTGGTGAA
AATGGCTCTCCTGGTGCCCTGGCGCTCCTGGTCTATCCAGGCCACCTGGTCTGTGGTCCAGCTGGAAAGAGT
GGTGACAGAGAGAAAGTGGCCCTGCTGGCCCTGCTGGTGCTCCCGGTCTGCTGGTTCCCGAGGTGCTCCTGGT
CCTCAAGGCCACGTGGTGACAAAGGTGAAACAGGTGAACGTGGAGCTGCTGGCATCAAAGGACATCGAGGATTC
CCTGGTAATCCAGGTGCCCCAGGTCTCCAGGCCCTGCTGGTTCAGCAGGGTGCAATCGGCAGTCCAGGACCTGCA
GGCCCCAGAGGACCTGTTGGACCCAGTGGACCTCCTGGCAAAGATGGAACCAAGTGGACATCCAGGTCCCATTGGA

988/6881
FIGURE 921B

CCACCAGGGCCTCGAGGTAACAGAGGTGAAAGAGGATCTGAGGGCTCCCCAGGCCACCCAGGGCAACCAGGCCCT
CCTGGACCTCCTGGTGCCCCCTGGTCCTTGCTGTGGTGGTGTGGAGCCGCTGCCATTGCTGGGATTGGAGGTGAA
AAAGCTGGCGGTTTTGCCCGTATTATGGAGATGAACCAATGGATTTCAAAATCAACACCGATGAGATTATGACT
TCACTCAAGTCTGTTAATGGACAAATAGAAAGCCTCATTAGTCCTGATGGTTCTCGTAAAAACCCGCTAGAAAC
TGCAGAGACCTGAAATTCTGCCATCCTGAACTCAAGAGTGGAGAATACTGGGTTGACCCTAACCAAGGATGCAAA
TTGGATGCTATCAAGGTATTCTGTAATATGGAACTGGGGAAACATGCATAAGTGCCAATCCTTTGAATGTTCCA
CGGAAACACTGGTGGACAGATTCTAGTGCTGAGAAGAAACACGTTTGGTTTGGAGAGTCCATGGATGGTGGTTTT
CAGTTTAGCTACGGCAATCCTGAACTTCCTGAAGATGTCCTTGATGTGCAGCTGGCATTCTTCGACTTCTCTCC
AGCCGAGCTTCCCAGAACATCACATATCACTGCAAAAATAGCATTGCATACATGGATCAGGCCAGTGGAAATGTA
AAGAAGGCCCTGAAGCTGATGGGGTCAAATGAAGGTGAATTCAAGGCTGAAGGAAATAGCAAATTCACCTACACA
GTTCTGGAGGATGGTTGCACGAAACACACTGGGGAATGGAGCAAAACAGTCTTTGAATATCGAACACGCAAGGCT
GTGAGACTACCTATTGTAGATATTGCACCCATGACATTGGTGGTCCTGATCAAGAATTGGTGTGGACGTTGGC
CCTGTTTGCTTTTTATAAACCAAACCTCTATCTGAAATCCCAACAAAAAAATTTAACTCCATATGTGTTCCTCTT
GTTCTAATCTTGTCAACCAGTGCAAGTGACCGACAAAATTCAGTTATTTATTTCCAAAATGTTTGGAAACAGTA
TAATTTGACAAAGAAAAATGATACTTCTCTTTTTTGTCTGTTCCACCAAATACAATTCAAATGCTTTTTGTTTTA
TTTTTTTACCAATTCCAATTTCAAATGTCTCAATGGTGCTATAATAAATAAACTTCAACACTCTTTATGATAAC
AACACTGTGTTATATTCTTTGAATCCTAGCCCATCTGCAGAGCAATGACTGTGCTCACCAGTAAAAGATAACCTT
TCTTTCTGAAATAGTCAAATACGAAATTAGAAAAGCCCTCCCTATTTTAACTACCTCAACTGGTCAGAAACACAG
ATTGTATTCTATGAGTCCCAGAAGATGAAAAAATTTTATACGTTGATAAAACTTATAAATTTCAATTGATTAATC
TCCTGGAAGATTGGTTTAAAAAGAAAAGTGTAATGCAAGAATTTAAAGAAATATTTTAAAGCCACAATTATTTT
AATATTGGATATCAACTGCTTGTAAGGTGCTCCTCTTTTTTCTTGTCATTGCTGGTCAAGATTACTAATATTTG
GGAAGGCTTTAAAGACGCATGTTATGGTGCTAATGTACTTTTCACTTTTAACTCTAGATCAGAATTGTTGACTTG
CATTGAGAACATAAATGCACAAATCTGTACATGTCTCCATCAGAAAGATTCAATTGGCATGCCACAGGGATTCT
CCTCCTTCATCCTGTAAAGGTCAACAATAAAAACCAAATATGGGGCTGCTTTTGTACACTAGCATAGAGAATG
TGTGAAATTTAACTTTGTAAGCTTGATGTGGTTGTTGATCTTTTTTTTTCTTACAGACACCCATAATAAATA
TCATATTAAATTC

989/6881

FIGURE 922

MMSFVQKGSWLLALLHPTIILAQQEAVEGGCSHLGQSYADRDVWKPEPCQICVCDSGSVLCDDIICDDQELDCP
NPEIPFGECCA VCPQPTAPTRPPNGQGPQGPKGDPGPPGIPGRNGDPGIPGQPGSPGSPGPPGICESCPTGPN
YSPQYDSYDVKSGVAVGGLAGYPGPAGPPGPPGPPGTSGHPGSPGSPGYQGPPEGQAGPSGPPGPPGAIGPSG
PAGKDGESGRPRGERGLPGPPGIKGPAGIPGFPGMKHGRGFDGRNGEKGETGAPGLKGENGLPGENGAPGPMG
PRGAPGERGRPLPGAAGARGNDGARGSDGQGPFPPTAGFPSPGAKGEVGPAGSPGSNGAPGQRGEPGPQG
HAGAQQPPGPPGINGSPGGKGEMGPAGIPGAPGLMGARGPPGPAGANGAPGLRGGAGEPGKNGAKGEPGPRGERG
EAGIPGVPGAKGEDGKDGSPEPGANGLPGAAGERGAPGFRGPAGPNGIPGEKGPAGERGAPGPAGPRGAAGEPG
RDGVPGGPGMRGMPGSPGGPGSDGKPGPPGSQGESGRPGPPGPGSPGRGQPGVMGFPGPKGNDGAPGKNGERGGPG
GPGPQGPFGKNGETGPPQGPPTGPGGDKGDTGPPGPQGLQGLPPTGGPPGENGKPGEPGPKGDAGAPGAPGGKG
DAGAPGERGPPGLAGAPGLRGGAGPPGPEGKGAAGPPGPPGAAGTPGLQGMPPGERGGLGSPGPKGDKGEPGGPG
ADGVPGKDGPRGPTGPIGPPGAGQPGDKGEGGAPGLPLGIAGPRGSPGERGETGPPGPAGFPAGQNGEPGGKG
ERGAPGEKGEPPGVPAGPPGGSGPAGPPGPQGVKGERGSPGGPGAAGFPAGRLPGPPGSNGNPGPPGPSGSPG
KDGPPGPAGNTGAPGSPGVSGPKGDAGQPGKEKSGPAQGGPPGAPGLGIAGITGARGLAGPPGMPGPRGSPGPQG
VKGESGKPGANGLSGERGPFGPQGLPGLAGTAGEPGRDGNPGSDGLPGRDGSPGGKGDRENGSPGAPGAPGHPG
PPGPVGPAGKSGDRGESGPAGPAGAPGPAGSRGAPGPQGPGRDKGETGERGAAGIKHGRGFPNGNPGAPGSPGPAG
QQCAIGSPGPAGPRGPVGPSPGPPKDGTSHPGPIGPPGPRGNRGERGSEGSPPGHPGQPGPPGPPGAPGPCCGGV
GAAAIAGIGGEKAGGFAPYYGDEPMDFKINTDEIMTSLKSVNGQIESLISPDSRKNPARNCRDLKFCHPELKSG
EYWVDPNQGCKLDAIKVFCNMETGETCISANPLNVRKHWTDSSAEKKHVWFGESMDGGFQFSYGNPELPEDVL
DVQLAFLRLSSRASQNTIYHCKNSIAYMDQASGNVKALKLMGSNEGEFKAEGNSKFTYTVLEDGCTKHTGEWS
KTVFEYRTRKAVRLPIVDIAPYDIGGPDQEFQVDPVGPVCF

990/6881
FIGURE 923

ACGCGTCCGAGCTGGCTCAGGGCGTCCGCTAGGCTCGGACGACCTGCTGAGCCTCCCAAACCGCTTCCATAAGGC
TTTGCCTTTCCAACCTCAGCTACAGTGTAGCTAAGTTTGGAAAGAAGGAAAAAGAAAAATCCCTGGGCCCCCTT
TCTTTTGTCTTTGCCAAAGTCGTCGTTGTAGTCTTTTGGCCAAGGCTGTTGTGTTTTAGAGGTGCTATCTCC
AGTTCCTTGCACTCCTGTTAACAAGCACCTCAGCGAGAGCAGCAGCAGCGATAGCAGCCGCAGAAGAGCCAGCGG
GGTCGCCTAGTGTCATGACCAGGGCGGGAGATCACAACCGCCAGAGAGGATGCTGTGGATCCTTGGCCGACTACC
TGACCTCTGCAAAATTCCTTCTCTACCTTGGTCATTCTCTCTACTTGGGGAGATCGGATGTGGCACTTTGCGG
TGTCIGTGTCTTCTGGTAGAGCTCTATGGAACAGCCTCCTTTTGACAGCAGTCTACGGGCTGGTGGTGGCAGGGT
CTGTTCTGGTCCTGGGAGCCATCATCGGTGACTGGGTGGACAAGAATGCTAGACTTAAAGTGGCCAGACCTCGC
TGGTGGTACAGAAIGTTTCAGTCATCCTGTGTGGAATCATCTGATGATGGTTTTCTTACATAAACATGAGCTTC
TGACCATGTACCATGGATGGGTTCTCACTTCTGTCTATATCTGATCATCACTATTGCAAAATTTGCAAAATTTGG
CCAGTACTGCTACTGCAATCACAATCCAAAGGGATTGGATTGTTGTTGTTGTCAGGAGAAGACAGAAGCAAACCTAG
CAAATATGAATGCCACAATACGAAGGATTGACCAGTTAACCAACATCTTAGCCCCATGGCTGTTGGCCAGATTA
TGACATTTGGCTCCCCAGTCATCGGCTGTGGCTTTATTTGCGGATGGAACCTGGTATCCATGTGCGTGGAGTACG
TCCTGCTCTGGAAGGTTTACCAGAAAACCCAGCTCTAGCTGTGAAAGCTGGTCTTAAAGAAGAGGAAACTGAAT
TGAAACAGCTGAATTTACACAAAGATACTGAGCCAAAACCCCTGGAGGGAACCTCATTAATGGGTGTGAAAGACT
CTAACATCCATGAGCTTGAACATGAGCAAGAGCCTACTTGTGCCTCCAGATGGCTGAGCCCTTCCGTACCTTCC
GAGATGGATGGGTCTCCTACTACAACCAGCCTGTGTTTCTGGCTGGCATGGGTCTTGCTTTCCTTTATATGACTG
TCCTGGGCTTTGACTGCATCACCACAGGGTACGCCCTACACTCAGGGACTGAGTGGTTCATCCTCAGTATTTGA
TGGGAGCATCAGCTATAACTGGAATAATGGGAACGTAGCTTTTACTTGGCTACGTCGAAAATGTGGTTTGGTTC
GGACAGGTCTGATCTCAGGATTGGCACAGCTTTCTGTTTGAICTTGTGTGTGATCTCTGTATTTCATGCCCTGGAA
GCCCCCTGGACTTGTCCGTTTCTCCTTTTGAAGATATCCGATCAAGGTTCAATTCAAGGAGAGTCAATTACACCTA
CCAAGATACCTGAAATTACAACGAAATATACATGTCTAATGGGTCTAATTCTGCTAATATTGTCCCGGAGACAA
GTCCTGAATCTGTGCCATAATCTCTGTCTGCTGCTGTTTGCAGGCGTCATTGCTGCTAGAATCGGTCTTTGGT
CCTTTGATTAACTGTGACACAGTTGCTGCAAGAAAATGTAATTGAATCTGAAAGAGGCATTATAAATGGTGTAC
AGAACTCCATGAAGTATCTTCTTGATCTTCTGCATTTTCATCATGGTCATCCTGGCTCCAAATCCTGAAGCTTTTG
GCTTGCTCGTATTGATTTAGTCTCCTTTGTGGCAATGGGCCACATTATGTATTTCCGATTTGCCCCAAATACTC
TGGGAAACAAGCTCTTTGCTTGGCGTCTGATGCAAAAGAAGTTAGGAAGGAAAATCAAGCAAATACATCTGTTG
TTTGAAGACAGTTTAACTGTTGCTATCCTGTTACTAGATTATATAGAGCACATGTGCTTATTTTGTACTGCAGAAT
TCCAATAAATGGCTGGGTGTTTTGCTCTGTTTTTACCACAGCTGTGCCTTGAGAACTAAAAGCTGTTTAGGAAAC
CTAAGTCAGCAGAAATTAAGTATTAATTTCCCTTATGTTGAGGCATGGAAAAAAATTTGAAAAAGAAAACTCA
GTTTAAATACGGAGACTATAATGATAACACTGAATTTCCCTATTTCTCATGAGTAGATACAATCTTACGTAAAAG
AGTGGTTAGTCACGTGAATTCAGTTATCATTGACAGATTCTTATCTGTACTAGAATTCAGATATGTCAGTTTTT
TGCAAACTCACTCTTGTTCAGACTAGCTAATTTATTTTTTGCATCTTAGTTATTTTTTAAAAACAAATTTCTC
AAGTATGAAGACTAAATTTTGATAACTAATATTATCCTTATTGATCCTATTGATCTTAAGGTATTTACATGTATG
TGGAAAAACAAACACTTAAGTAGAATCTCTAATAAGGTTTATGGTTTAGCTTAAAGAGCACCTTTGTATTTTT
ATTATCAGATGGGGCAACATATTGTATGAAGCATATGTAGCACTTCACAGCATGGTTATCATGTAAGCTGCAGGT
AGAAGCAAAGCTGTAAAGTAGATTTATCACACAATGACTGCATACAGACTTCAAATATGTCAATAGTTTGGTCAT
AGAACCTAGAAGCCAAAAGCCACACAGAAGGGCAAGAATCCCAATTTAACTCATGTTATCATCATTAGTGATCTG
TGTGTAGAACATGAGGGTGTAAAGCCTTCAGCCTGGCAAGTTACATGTAGAAAGCCACACTTGTGAAGTTTTG
TTTTACAAATCACTTGATTTAACACACTCAGGTAGAATATTTTTTATTTTACTGTTTTATACCCAGAAGTTATTT
CTACATTGTTCTACAGCAAGAATATTCATAAAAGTATCCCTTTCAAATGCCCTTGAGAAGAATAGAAGAAAAAA
GTTTGTATATATTTTTAAAAAATTGTTTTAAAAGTCAGTTTGAACATGTCTGTACCAAGATGGTACTTTGCCTTA
ACCGTTTATATGCATTTTCATGGAGACTGCAATACGTTGCTATGAGCACTTTCTTTATCCTTGGAGTTTAACTCT
TTGCTTCATCTTTCTACAGTATGACATAATGATTTGCTATGTTGTAAAATCTTTGTAAAAAATTTCTATATAAAA
ATATTTTGAAGTCTTAAAAAATTTTTTTTTT

991/6881
FIGURE 924

MTRAGDHNQRGCCGSLADYLTSAKFLLYLGHSLSTWGDRMWHFAVSVFLVELYGNLLLLTAVYGLVVAGSVLVL
GAIIGDWVDKNARLKVAQTSLVVQNVSVILCGIILMMVFLHKHELLTMYHGWLTSCYILIITIANIANLASTAT
AITIQRDWIVVVAGEDRSKLANMNATIRRIDQLTNILAPMAVGQIMTFGSPVIGCGFISGWNLVSMCVEYVLLWK
VYQKTPALAVKAGLKEEETELKQLNLHKDTEPKPLEGTHLMGVKDSNIHELEHEQEPTCASQMAEPFRTRDGVV
SYYNQPVFLAGMGLAFLYMTVLGFDCITTGYAYTQGLSGSILSILMGASAITGIMGTVAFTWLRRKCGLVRTGLI
SGLAQLSCLILCVISVFMPGSPDLDSVSPFEDIRSRFIQGESITPTKIPETITEIYMSNGSNSANIVPETSPEV
PIISVSLLFAGVIAARIGLWSFDLTVTQLLQENVIESERGIINGVQNSMNYLLDLLHFIMVILAPNPEAFGLLV
ISVSFVAMGHIMYFRFAQNTLGNKLFACGPDAKEVRKENQANTSVV

992/6881
FIGURE 925

TTTTGTGAAGAGACGAAGACTGAGCGGTTGCGGCCGCGTTGCCGACCTCCAGCAGCAGTCGGCTTCTCTACGCAG
AACCCGGGAGTAGGAGACTCAGAATCGAATCTCTCTCCCTCCCCTTCTTGTTTTCGGCTTTGTGAGAAACCTTA
CCATCAAAACACAATGGCCAGCAACGTTACCAACAAGACAGATCCTCGTTCCATGAACTCCCGTGTATTTCATTGGG
AATCTCAACACTCTTGTTGGTCAAGAAATCTGATGTGGAGGCAATCTTTTCGAAGTATGGCAAAATTGTGGGCTGC
TCTGTTTCATAAGGGCTTTGCCTTCTTTTCAGTATGTTAATGAGAGAAATGCCCGGGCTGCTGTAGCAGGAGAGGAT
GGCAGAATGATTGCTGGCCAGGTTTTAGATATTAACCTGGCTGCAGAGCCAAAAGTGAACCGAGGAAAAGCAGGT
GTGAAACGATCTGCAGCGGAGATGTACGGCTCCTCTTTGACTTGGATTGTGACTTCAACGGGACTATTATGAT
AGGATGTACAGTTACCCAGCACGTGTACCTCCTCCTCCTATTGCTCGGGCTGTAGTGCCTCGAAACGTCAG
CGTGTATCAGGAAACACTTCACAAAGGGGCATAAGTGGCTTCAATTCTAAGAGTGGACAGCGGGGATCTTCCAAG
TCTGGAAAGTTGAAAGGAGATGACCTTCAGGCCATTAAGAGGGAGCTGACCCAGATAAAAACAAAAGTGGATTCT
TTCCTGGAAAACCTGGAAAAAATTGAAAAGGAACAGAGCAAAACAGCAGTAGAGATGAACAATGTTAAGTCAGAA
GAGGAGCAGAGCAGCAGCTCCGTGAAGAAAGATGAGACTAATGTGAAGATGGAGTCTGAGGGGGGTGCAGATGAC
TCTGCTGAGGAGGGGGACCTACTGGATGATGATGATAATGAAGATGGGGGGATGACCAGCTGGAGTTGATCAAGG
ATGATGAAAAAGAGGCTGAGGAAGGAGAGGATGACAGAGACAGCGCCAATGGCGAGGATGACTCTTAAGCACATA
GTGGGGTTTTAGAAATCTTATCCATTATTTCTTTACCTAGGCGCTTGTCTAAGATCAAATTTTTACCAGATCCT
CTCCCCTAGTATCTTCAGCACATGCTCACIGTTTTCCCCATCCTTGTCTTCCCATGTTTCAATTAATTAATTCATA
TTGCCCCGCGCCTAGTCCCATTTTCACTTCCTTTGACGCTCCTAGTAGTTTTGTTAAGTCTTACCCTGTAATTTT
TGCTTTTAAATTTTGATACCTCTTTATGACTTAACAATAAAATGATGTATGGTTTTTATCAACTGTCTCCAAAAT
AATCTCTTGTATGCAGGGAGTACAGTTCTTTTCATTACATACATAAGTTTCAGTAGTTGCTTCCCTAACTGCAAAG
GCAATCTCATTTAGTTGAGTAGCTTTGAAAGCAGCTTTGAGTTAGAAGTATGTGTGTTACACCCCCACATTAGT
GTGCTGTGTGGGGCAGTTCAACACAAATGTAACAATGTATTTTTGTGAATGAGAGTTGGCATGTCAAATGCATCC
TCTAGAAAAATAATTAGTGTTATAGTCTTAAGATTTGTTTTCTAAAGTTGATACTGTGGGTATTTTTGTGAACA
GCCTGATGTTTGGGACCTTTTTTCTCAAAATAACAAGTCCTTATTAAACCAGG

993/6881
FIGURE 926

GCCGACCTCCAGCAGCAGTCGGCTTCTCTACGCAGAACCCGGGAGTAGGAGACTCAGAATCGAATCTCTTCTCCC
TCCCTTCTTGTTTTCGGCTTTGTGAGAAACCTTACCATCAAACACAATGGCCAGCAACGTTACCAACAAGACAG
ATCCTCGTTCCATGAACTCCCGTGTATTCAATTGGGAATCTCAACACTCTTGTGGTCAAGAAATCTGATGTGGAGG
CAATCTTTTCGAAGTATGGCAAAATTGTGGGCTGCTCTGTTTCATAAGGGCTTTGCCTTCTTTCAGTATGTTAATG
AGAGAAATGCCCGGGCTGCTGTAGCAGGAGAGGATGGCAGAATGATTGCTGGCCAGGTTTTAGATATTAACCTGG
CTGCAGAGCCAAAAGTGAACCGAGGAAAAGCAGGTGTGAAACGATCTGCAGCGGAGATGACCTTCAGGCCATTAA
GAGGGAGCTGACCCAGATAAAACAAAAGTGGATTCTTTCCTGGAAAACCTGGAAAAAATTGAAAAGGAACAGAG
CAAACAAGCAGTAGAGATGAACAATGTTAAGTCAGAAGAGGAGCAGAGCAGCAGCTCCGTGAAGAAAGATGAGAC
TAATGTGAAGATGGAGTCTGAGGGGGGTGCAGATGACTCTGCTGAGGAGGGGGACCTACTGGATGATGATGATAA
TGAAGATGGGGGGATGACCAGCTGGAGTTGATCAAGGATGAGGATGACTCTTAAGCACATAGTGGGGTTTAGAAA
TCTTATCCCATTTATTTCTTTACCTAGGCGCTTGTCTAAGATCAAATTTTTCACCAGATCCTCTCCCCTAGTATCT
TCAGCACATGCTCACTGTTTTCCCCATCCTTGTCTTCCCATGTTTATTAAATTAATTCATATTGCCCCGCGCCTA
GTCCCATTTTCACTTCCTTTGACGCTCCTAGTAGTTTTGTAAAGT

994/6881
FIGURE 927

GAGACGAAGACTGAGCGGTTGCGGCCGCGTTGCCGACCTCCCTTCTTGTTTTCGGCTTTGTGAGAAACCTTACC
ATCAAACACAATGGCCAGCAACGTTACCAACAAGACAGATCCTCGTTCCATGAACTCCCGTGATTTCATTGGGAA
TCTCAACACTCTTGTTGGTCAAGAAATCTGATGTGGAGGCAATCTTTTCGAAGTATGGCAAAATTGTGGGCTGCTC
TGTTTATAAGGGCTTTGCCTTCTTTCAGTATGTTAATGAGAGAAATGCCCCGGGCTGCTGTAGCAGGAGAGGATGG
CAGAATGATTGCTGGCCAGGTTTTAGATATTAACCTGGCTGCAGAGCCAAAAGTGAACCGAGGAAAAGCAGGTGT
GAAACGATCTGCAGCGGAGATGTACGGCTCCTCTTTTGAAGTTGGATTGTGACTTTCAACGGGACTATTATGATAG
GATGTACAGTTACCCAGCACGTGTACCTCCTCCTCCTCTATTGCTCGGGCTGTAGTGCCCTCGAAACGTCAGCG
TGTATCAGGAAACACTTCACAAAGGGGCATAAGTGCTTCAATTCTAAGAGTGGACAGCGGGGATCTTCCAAGTC
TGGAAGTTGAAAGGAGATGACCTTCAGGCCATTAAGAGGGAGCTGACCCAGATAAAACAAAAAGTGGATTCTTT
CCTGGAAAACCTGAAAAAATTGAAAAGGAACAGAGCAAACAAGCACTGGAGTTGATCAAGGATGATGAAAAAGA
GGCTGAGGAAGGAGAGGATGACAGAGACAGCGCCAATGGCGAGGATGACTCTTAAGCACATAGTGGGGTTTAGAA
ATCTTATCCCATTTATTTCTTTACCTAGGCGCTTGCTAAGATCAAATTTTTCACCAGATCCTCTCCCCTAGTATC
TTCAGCACATGCTCACTGTTTTCCCATCCTTGCTCCTCCCATGTTTCAATTAATTCATATTGCCCGCGCCT
AGTCCCATTTTCACTTCCTTTGACGCTCCTAGTAGTTTT

995/6881
FIGURE 928

ATCTTTTAGAGAAATCTCGGGTTGTTAAACAGCCAAGAGGTGAAAGAACTTCCATGTGTTCTATCAGCTGCTCT
CTGGTGCCTCTGAAGAGCTCCTCAATAAACTTAAGCTTGAGAGGGATTTCAGCAGGTATAACTACCTGAGTCTGG
ATTTCGGCCAAAGTGAATGGAGTGGATGATGCAGCAAATTTAGAACCGTGCAGGAATGCCATGCAGATTGTGGGCT
TTATGGATCATGAAGCTGAGTCTGTCTTGGCGGTGGTGGCAGCAGTGTGAACTGGGGAAACATTGAGTTCAAGC
CCGAATCTCGAGTGAATGGTCTAGATGAAAGCAAATCAAAGATAAAAAATGAGTTAAAAGAAATTTGTGAATTGA
CCGCATTGATCAATCAGTTCTAGAACGAGCATTTCAGTTTCCGAACAGTTGAGGCCAAACAGGAGAAAGTTTCAA
CTACACTGAATGTGGCTCAGGCTTATTATGCCCCTGATGCTCTGGCTAAAAACCTCTACAGCAGGTGTGTTTTCAT
GGTTGGTAAATCGAATCAATGAAAGCATTAAAGGCACAAACAAAAGTGAGAAAGAAGTTCATGGGTGTTCTGGACA
TTTATGGCTTTGAGATTTTCGAGGACACAGCTTTGAGCAGTTTATTATTAATTATTGTACGAAAAGCTGCAAC
AAATCTTCATTGAACTTACTCTTAAAGAAGAGCAGGAGGAGTATATACGGGAGGATATAGAATGGACTCACATTG
ACTACTTCAATAATGCTATCATTTGTGACCTAATAGAAAATAACACAAATGGAATCCTGGCCATGCTGGATGAAG
AGTGCCCTCAGACCTGGCACAGTCACTGATGAGACCTTCTTAGAAAAGCTGAACCAAGTATGTGCCACCCACCAGC
ATTTTGAGAGCAGGATGAGCAAGTGCTCTCGGTTCTCTCAATGACACGCTCTCTGCCTCACAGCTGCTTCAGGATCC
AGCATTATGCTGGAAGGTGCTGTACCAGGTGGAAGGATTTCGTTGACAAAAACAATGACCTTCTCTATCGAGACC
TGTCCTCAAGCCATGTGGAAGGCCAGCCATGCCCTCATCAAGTCTTTGTTCCCCGAAGGGAATCCCGCCAAGATCA
ACCTGAAAAGGCCTCCTACAGCAGGCTCACAGTTCAAGGCATCCGTGGCCACTCTGATGAAAACCTACAGACCA
AGAACCCAAACTATATTAGGTGTATCAAACCGAATGATAAAAAAGCAGCACACATCTTCAACGAGGCTCTAGTGT
GTCATCAGATCAGGTACCTGGGGCTTTTGGAGAACGTCAGTGCAGGAGGAGGCTACGCCTTCAGGCAGGCCCT
ATGAACCTTGCCCTAGAAAAGATACAAAATGCTTTGTAAACAAACATGGCCCTCATTTGGAAGGACCAGCCAGGTCTG
GTGTGGAGGTCTTATTAATGAATTAGAAATTCCTGTTGGAAGAATACTCCTTTGGTAGATCAAAGATATTTCATCC
GAAACCCAAGAACATTATTCAAATTAGAAGACCTGAGGAAGCAACGCCTGGAGGACTTGGCCACTCTCATTGAGA
AGATATATCGGGGTGGAAATGCCGCACACACTTCTGTCTAATGAAAAAAGCCAAATTGTGATTGCCGCCTGGT
ACAGGAGATATGCGCAACAAAAGAGGTACCAGCAGACAAAGAGTTCCGCCTTAGTAATTCAGTCTTATATCCGGG
GTTGGAAGGCTCGAAAAATTCTGCGGAACTGAAGCATCAAAAGCGCTGTAAGGAAGCAGTCACGACCATTGCTG
CATATTGGCATGGGACCCAGGCTCGAAGGGAATTGAAACGCTTGAAGGAGGAGGCTAGGCGTAAGCATGCAGTTG
CTGTCATTTGGGCTTACTGGCTTGGACTGAAGGTACGTAGAGAATACAGGAAATCTTCAGAGCCAATGCTGGAA
AGAAAATCTATGAGTTTACGCTTCAGAGAATTGTGCAAAAATACTTCTTGGAAATGAAAAATAAGATGCCTTCCT
TATCTCCAATAGACAAGAATTGGCCCTCAAGACCTTACTTATCTTGGATTCTACTCACAAGGAGCTAAAAAGGA
TTTTCCACTTGTGGAGGTGTA AAAAATACAGGGACCAATTCACAGACCAGCAGAACTTATTTATGAAGAGAAAC
TAGAAGCCAGTGAACCTTCAAAGACAAGAAGGCTTTATACCCATCTAGTGTGTTGGGCAACCATTCCAAGGGGCTT
ACCTGGAAATCAACAAGAACCCCAAGTATAAGAACTCAAAGATGCCATTGAAGAAAAGATCATATTGCTGAAG
TCGTGAACAAAATTAACCGTGCTAATGGGAAGAGTACATCTCGGATTTTCTCTTAACAAACAATAATCTCCTTC
TTGCTGACCAAAAGTCTGGACAAATCAAGTCAGAGGTTCCATTGGTGGATGTGACCAAGGTATCAATGAGCTCAC
AAAATGATGGCTTCTTCGCCGTCCACCTCAAAGAGGGCTCAGAAGCAGCTAGTAAAGGAGACTTCTCTTCAGCA
GTGATCACCTGATTGAAATGGCCACCAAGCTCTATCGCACAACTCTCAGCCAAACCAACAGAAGCTCAATATTG
AGATTTCCGATGAGTTCTGTTACAGTTTACAGCAGGACAAAGTATGTGTGAAGTTTATTAGGAACCTGTAGTAT
GATATTTAACAATATAGGCTTCAAGAAGGGCTGCTCCTAAGAGGGGCGAGAAATGAATGACCAGGTAAATCCCT
CTACATGTGGTTTCTGTTTG

996/6881
FIGURE 929

AGGTCAGAAATGTGAATTCCAGGATGCCTATGTTCTGTTGAGTGAAAAGAAAATTTCTAGTATCCAGTCCATTGT
ACCTGCTCTTGAAATTGCCAATGCTCACCGTAAGCCTTTGGTCATAATCGCTGAAGATGTTGATGGAGAAGCTCT
AAGTACACTCGTCTTGAATAGGCTAAAGGTTGGTCTTCAGGTTGTGGCAGTCAAGGCTCCAGGGTTTGGTGACAA
TAGAAAGAACCAGCTTAAAGATATGGCTATTGCTACTGGTGGTGCAGTGTTTGGAGAAGAGGGATTGACCCGTGA
TCTTGAAGACGTTTCAGCCTCATGACTTAGGAAAAGTTGGAGAGGTCATTGTGACCAAAGACGATGCCATGCTCTT
AAAAGGAAAAGGTGACAAGGCTCAAATTGAAAAACGTATTCAAGAAATCATTGAGCAGTTAGATGTCACAACTAG
TGAATATGAAAAGGAAAACTGAATGAACGGCTTGCAAACTTTTCAGATGGAGTGGCTGTGCTGAAGGTTGGTGG
GACAAGTGATGTTGAAGTGAATGAAAAGAAAGACAGAGTTACAGATGCCCTTAATGCTACAAGAGCTGCTGTTGA
AGAAGGCATTGTTTTGGGAGGGGGTTGTGCCCTCCTTCGATGCATTCCAGCCTTGGACTCATTGACTCCAGCTAA
TGAAGATCAAAAAATTGGTATAGAAATTATTAAAAGAACTCAAAATTCAGCAATGACCATTGCTAAGAATGC
AGGTGTTGAAGGATCTTTGATAGTTGAGAAAATTATGCAAAGTTCCCTCAGAAAGTTGGTTATGATGCTATGGCTGG
AGATTTTGTGAATATGGTGGAAAAAGGAATCATTGACCCAACAAAGGTTGTGAGAACTGCTTTATTGGATGCTGC
TGGTGTGGCCTCTCTGTAACTACAGCAGAAGTTGTAGTCACAGAAATTCCTAAAGAAGAGAAGGACCCTGGAAT
GGGTGCAATGGGTGGAATGGGAGGTGGTATGGGAGGTGGCATGTTCTAACTCCTAGACTAGTGCTTTACCTTTAT
TAATGAACTGTGACAGGAAGCCCAAGGCAGTGTTCCCTACCAATAACTTCAGAGAAGTCAGTTGGAGAAAATGAA
GAAAAAGGCTGGCTGAAAATCACTATAACCATCAGTTACTGGTTTCAGTTGACAAAATATATAATGGTTTACTGC
TGTCATTGTCCATGCCTACAGATAATTTATTTTGTATTTTGAATAAAAAACATTTGTACATTCCCTGATACTGGG
TACAAGAGCCATGTACCAAGTGTACTGCTTTCAACTTAAATCACTGAGGCATTTTACTACTATTCTGTTAAATC
AGGATTTTAGTGCTTGCCACCACCAGATGAGAAGTTAAGCAGCCTTTCTGTGGAGAGTGAGAATAATTGTGTACA
AAGTAGAGAAGTATCCAATTATGTGACAACCTTTGTGTAATAAAAAATTTGTTTAA

997/6881

FIGURE 930

GCTACACTAGAGCAGAGTACGAGTCTGAGGCGGAGGGAGTAATGGCAGGACAAGCGTTTAGAAAGTTTCTTCCAC
TCTTTGACCGAGTATTGGTTGAAAGGAGTGCTGCTGAAACTGTAAACCAAAGGAGGCATTATGCTTCCAGAAAAAT
CTCAAGGAAAAGTATTGCAAGCAACAGTAGTCGCTGTTGGATCGGGTTCTAAAGGAAAGGGTGGAGAGATTCAAC
CAGTTAGCGTGAAAGTTGGAGATAAAGTTCTTCTCCAGAATATGGAGGCACCAAAGTAGTTCTAGATGACAAGG
ATTATTTCCCTATTAGAGATGGTGACATTCTTGGAAGTACGTAGACTGAAATAAGTCACTATTGAAATGGCATC
AACATGATGCTGCCCATTCCTGAAGTTCTGAAATCTTTCGTCATGTAAATAATTCCATATTTCTCTTTATA
ATAAACTAATGATAACTAATGACATCCAGTGTCTCCAAAATTGTTTCCTTGTAAGTATATAAACTTCCAAATA
AAAAATATGTAAAT

998/6881
FIGURE 931

MAGQAFRKFLPLFDRVLVERSAAETVTKGGIMLPEKSQGKVLQATVVAVGSGSKGKGGEIQPVSVKVGDKVLLPE
YGGTKVVLDDKDYFLFRDGDILGKYVD

999/6881
FIGURE 932

GATTTGACCCTTGAGCCGTAGGGAGCGCGGCATTTTCTGGAAAGTTCTGGAACCGAGCGAGGCCCGGGAAC TAGA
CTAAGCCGGCCGGAGAGGGCTGAGCGCGCTAGCACACCCTGCGCGGAAATGCTTCGGTTACCCACAGTCTTTCGC
CAGATGAGACCGGTGTCCAGGGTACTGGCTCCTCATCTCACTCGGGCTTATGCCAAAGATGTAAATTTGGTGCA
GATGCCCCGAGCCTTAATGCTTCAAGGTGTAGACCTTTTAGCCGATGCTGTGGCCGTTACAATGGGGCCAAAGGGA
AGAACAGTGATTATTGAGCAGAGTTGGGGAAGTCCCAAAGTAACAAAAGATGGTGTGACTGTTGCAAAGTCAATT
GACTTAAAAGATAAATACAAAACATTGGAGCTAAACTTGTTCAAGATGTTGCCAATAACACAAATGAAGAAGCT
GGGGATGGCACTACCACTGCTACTGTACTGGCACGCTCTATAGCCAAGGAAGGCTTCGAGAAGATTAGCAAAGGT
GCTAATCCAGTGGAATCAGGAGAGGTAGGAATGTCTGTTGTCACCACAGTGTTCTAAACTTCAGTGTGCTTTAA
GGCATTGATTCCCTTGACCTAAGGAAAGTGTTAACTTGAGGAAATTAATCCTTCTCTCATTCTCTCTTGGCCT
GGAGAAATCTTAAACTGGGTTTTCTTTAGCAAGCTTGGCTGT

1000/6881
FIGURE 933

TGGAAAGAGTGGAAACGAGAAAAACCTTTTCAGACTATTGTGTTCTGGGCCAGCGTCCAATGCATTTACCAAATATGA
ACCAGCTGGCATCCCTGGGGAAAACCAACGAACAGTCTCCTCAAGCCAAATTCACCACAGTACTCCAATCCGAA
ACCAAGTGCCCGCATTACAGCCCATCATGAGCCCTGGTCTTCTTTCTCCCAGCTTAGTCCACAACCTTGTAAGGC
AACAAATAGCCATGGCCCCATCTGATAAACCAACAGATTGCCGTTAGCCGGCTCCTGGCTCACCAGCATCCTCAAG
CCATCAACCAGCAGTTCCTGAACCATCCACCCATCCCAGAGCAGTTAAGCCAGAGCCAACCAACTCTTCCGTGG
AAGTCTCTCCAGATATCTACCAGCAAGTCAGAGATGAGCTGAAGAGGGCCAGTGTGTCCCAAGCTGTCTTTGCAA
GAGTGGCATTCAACCGCACACAGGTACAATTAGCATTAAACACTGTAATTAACAGTAATACTGGGGACAGAATTG
AGATAGGTTATAATTATTTTATCTCTTTAAAAAGCTTTATGGATCTCAGGACATAGAATTAGATTAAATTAATA
TCCTTAGTTAATGAAACGCTATTACCATTGATCCATGCATGACCATGATTAAATTAATAATTGATTAAATTTGAT
GAGGTAATGGACATCTTTGAACCCAACATGCAACTCAGGACCCCAATGTTACTAATGACTTGGGTATGTACTCT
CTCATCTCTCCATTTACTGTTTGCAGATGTAACCTGATCTTGAATTTTGTGATTATCAGTCTCTTGCCATTTT
TTAAAACAAATTATAAACTATGCATGCCAGACAATCTATTTTTTCCCTTGGTTTTGAACGTTATAAAAGAAA
TAAATTATGTTTTATAATTTTATAAGTAAAGTTTTATTTAAAAATTTATAAAAATCCTTCAAAACATAAAAGTT
AGATTTAAGATATTGATAAAATGATAATGGATAGTTTACTTTTATTTTTAATGTTTGAGAGAATTATATGCCAAT
TATAACACTAAATGTTGAAAAATGTAAACAGTTAATTCACACGCTCTATTTTTTATTGAGGGTGCTTATATGTAAGA
CATTCTTTTCTGCTAGTAGTTCTTAAACTGGAGTGTGCATCAGAATCACCTGGAGGGCTGTGAAACACAGACTGC
TGGGCCTCATCCCATAGTTTCTGATTGAGTAAGTCTGAGGTTGGACTCAGAAATGGACATTTCTAACAAGATCCC
AGGTGATTATGATGGTACTGATCCGGGGCCGCATGTTGAGAAAACTTACCTAGCCCTTATATGAGTAAGAACA
AGCAGATGTGCATAATCTCGGGGAGCTAACGATCTAGTGGTGAAGAACAGGTAACCAGATAATTAGAGTAAATG
TGAAGTGTTATGAGTAATGTTGGAGAGTGTAAAGCAATAGTTGTGGTGGAGGTTTCATGGTGAGCTGGAGGTGAGGA
ATACTTCCAGGAGAATGAATATGGTAAGCCTTCAAGAAAACAGTAGATTAAATTTTCAGCACCCCTCGTAAATGCA
AGGCAGGAAGCTAAGCTCTATGTGGGTGCAAAGAGGAAAAAGATAATGGGCAAGCACATATGTAAACAACACTCT
AGAGTATTAGGTTGTCAATGGGTTAGACCTTGAAGGATTAATAAAGTACAGCAGGCAAGATGATGAAGAAGAAG
ATTATTATTTTAAGGTAAAGACATTCATAGGAAAGATGTTGAGTTAAAAATGCAGGAATAATTTGAAGACATAA
AACAACTGATTTCTGAGAAAAACCTGTGAACGTACAGAATGTAGGAATGCTTTTATTATCTGATACCTGTTCA
GAGACACATGATAACACACGCTGTAGATGGACCTTACAAATGAAAGAAAGCATGCTGGATTGAAACCTCAATAAT
TTGGAATAACAGGAATGTTTCCAAATTTAATGATTACTTCAGAAGTTAAGGGGCTGGGCACAGTGGCTCATGTC
TGTAAGTCTAGCACTTTGGGAGGCCAAGGTGGGCGGATCACTTGAGGTGAGGAGTTCAAGACCAGCCTGGGCAAG
ATGGTAAACACCATCTCTACTAAAACAGAACAAAAACAAAAAATTAGCTGGGCGTGGTGATGCGTGCCCTGTAGT
CTCAGCTACTCAGGAGGCTGAGGCAGAAATCCCTTGAACCCAGGAGGCGGAGGCTGCAGTGAGCCGAGATCAT
GCCACTGCACTCCAGCCTGAGTGAGAGTGAACTCCATCTCAAAAAAAAAAAAAAAAAA

1001/6881
FIGURE 934

GAGAGGAGACACCGCCGAGTTGCCGGTACATCGGGGATTTCTGGCTCTTTCCCTCTTCGCCCTTAAATTCGGGTGT
CTTTTATGATAATCAAAAGCAGCAAAGCCAAGCTATCAGGCCAGCGTTTTAAAACTAGAAAAAGAGATGAAA
AAGAGAGGTTTGACCTTACTCAGTTTCAAGACTGTATTATTCAAGGCTTAACTGAAACCGGTACTGATTTGGAAG
CAGTAGCTAAGTTTCTTGATGCTTCTGGAGCAAACTTGATTACCGTCGATATGCAGAAACACTCTTTGACATTC
TGGTGGCTGGTGAATGCTGGCCCCAGGTGGTACACTGGCAGATGACATGATGCGTACAGATGCTCTGCGTGTG
CAGCCCAAGAAGATCTAGAGACCATGCAAGCATTGCTCAGGTTTTTAAACAAGTTAATCAGGCGCTACAAATACC
TGGAGAAAGGTTTTGAAGATGAAGTAAAAAAGCTGCTGCTGTTCTTGAAGGGTTTTTCAGAGTCGGAGAGGAACA
AGCTAGCTATGTTGACTGGTGTTCTTCTGGCTAATGGAACACTTAATGCATCCATTCTTAATAGCCTTTATAATG
AAAAATTGGTTAAAGAAGGAGTTTTCAGCAGCTTTTGTCTGTGAAGCTCTTTAAATCATGGATAAATGAAAAAGATA
TCAATGCAGTAGCTGCAAGTCTTCGGAAAGTCAGCATGGATAACAGACTGATGGAACCTTTTCTGCCAATAAGC
AAAGTGTTGAACACTTCACAAAATATTTTACTGAGGCAGGCTTGAAAGAGCTTTTCAGAAATGTTTCGGAATCAGC
AAACCATCGGAGCTCGTAAGGAGCTCCAGAAAGAAGCTTCAAGAACAGATGTCCCGTGGTGATCCATTTAAGGATA
TAATTTTATATGTCAAGGAGGAGATGAAAAAAACAACATCCCAGAGCCAGTTGTCATCGGAATAGTCTGGTCAA
GTGTAATGAGCACTGTGGAATGGAACAAAAAGAGGAGCTTGTAGCAGAGCAAGCCATCAAGCACTTGAAGCAAT
ACAGCCCTCTACTTGCTGCCTTTACTACTCAAGGTGAGTCTGAGCTGACTCTGTTACTGAAGATTTCAGGAGTATT
GCTATGACAACATTCATTTTCATGAAAGCCTTCAGAAAAATAGTGGTGCCTTTTTTATAAAGCTGAAGTCCTGAGCG
AGGAGCCCATTTTGAAGTGGTATAAAGATGCACATGTTGCAAAAGGGGAGAGAGTGTTTTCTTGAGCAAAATGAAAA
AGTTTGTAAGTGGCTCAAAAATGCTGAAGAAGAATCTGAATCTGAAGCTGAAGAAGGTGACTGAATTTTGAAAC
TACACCCTCAGTAAAGCAAACAGGAGTTGTAGATAAAATGTCATGTCTCATGTGTCTGTTCTTACATCTTCTCT
ACCTCCCTGTATCAAGCATGATATAAGGGCTTTCATGGCAAATTTTATTTTAACTGTTTCTATGGTTGCTGGAAA
TGTTGGGTTTAGTTTTCTAAACCATGTTTTAAGTAGCTACAGGAGCTATAGATTTGAATCTAATGTTGCATTAGT
CTTTTCAGTTATCTTCTACCTCCTGTATTTTCTACTGTAATAATGTAATTTAAGGCCTTCCACAATGAACAGTTC
ACTTTATTCCTGGGTTTTCTATAAACAGTTTTAAGGATATGATTTGGTTAAAAAATAATTTGTTATAAAAAATTC
TGTTTGCAAATTAACCTGGAAAAGTATCCAGAGTCTCAAAGGCAATGATTTGTGAGATAATATGGCATGCCCGG
AGCCCTGCTCATCAATGAAAAACCCATATGTAATAATCGAATTCATTTAACATGAATCTTGAGTACGTGGACCAT
TGCTTGCGATGTTAACTTTTTGTTTTGTTTTGTTTTGTTTTGTTTTGCAATTTTAACTCCAGATATCCTAAAGCTC
AATTGTTTGGTCTCTGGTTTTTCATCCTTAGAGAAGCCATGGAGAACAGACTTGAAAAGTTTAGGAAATCATAATG
TGGCAGAGGTGGTGGGAAGAAGAAAGTTGAGCTTTTTCCCTTGAGAACTTCTGCATTTAGTTTCTATCTTTCC
AGGCAAAACAAATGGGTATCTTTTCATACAACCATTTTCAAATGAACCTTAGAAAAGCTTTAACATTTAAGGTA
TTTTATGCACAGAATACACTTAGATTGATAGGAAAGAAGCTGTAATGGAGTTTGAGTAAAGAAAAATGACTGATGT
ACTAAACCCAGTAAAAATTTGTTGAAAATGTTAAAGGTCAGCATGTTCTAATTGGGAATCTAGATATAGCTTAGAT
TTCTATTGGCTTAGAGTATTGCTATACCAAAATGAAGTGCATGACAATTATATATTCTACTCGGTCTACTG
GACTGGCTTCGTTCTCTTAATACTACTCAGTAATGACTCAAGCCTCTGGCTATTAAACATACCCTAGTTGCCGTTTT
TTAATTGCCATGACCCAAATACTTCTTGGTATACAATTGATCCATTTATTTTAAATGGCTGCCTTTTTCATTTTCAT
CTTTTCTGCTGCTACCCATCTATGTATGTAGTCAATTGGGGGGAAAAATGTAGCCACATTTTTTATGGGAAGACTT
TGTTGTTAAAGTGAACATTTTGAAGGTTTTTAACTGGTGAAACTAGCCTGGAATAATGCCACCAGAGACTGAGTG
GAAATCGCCCCCTTTTGAAGGTGCCATTCTTATGAGCCAAAAGTTTGTCATTTAAAGGTTTCAATTTGAGGGAATAA
CATGTAATATAAATTTGAAATAAAGGTATAGTAACCTTAAAAAGAACATTATAACTGATTGTTGTGAATGGGGTGA
ATTTGTTAAATGAGTAACCTTGATAAAGTTTTTCATGCACAGGCAAAATGTATTCAGTAGATTTCTACGTAGTG
ATCTGCTTTTACTTTGTAATTTGTAGTTCTCAAAAGACTTTTTTTTAAAAAATAAAGTCCATACCTTACACTT

1002/6881
FIGURE 935

MNNQKQKPTLSGQRFKTRKRDEKERFDPTQFQDCIIQGLTETGTDLEAVAKFLDASGAKLDYRRYAETLFDILV
AGGMLAPGGTLADDMMRTDVCVFAAQEDLETMQAFAQVFNKLIRRYKYLEKGFEDVKKLLFLKGFSESERNKL
AMLTGVLLANGTLNASILNSLYNENLVKEGVSAFAVKLFKSWINEKDINAVAASLRKVSMDNRLMELFPANKQS
VEHFTKYFTEAGLKELSEYVRNQQTIGARKELQKELQEQMSRGDPFKDIILYVKEEMKKNNIPEPVVIGIVWSSV
MSTVEWNKKEELVAEQAIKHLKQYSPLLAFTTQGQSELTILLKIQEYCYDNIHFMKAFQKIVVLFYKAEVLSEE
PILKWKDAHVAKGKSVFLEQMKKFVEWLKNAEESESEAEEGD

1003/6881
FIGURE 936

ACGCAGAGAAGTTTCCGGGACTGAAACTTGAACCTTTACCTGATTTCTGTATGTTGTCATCTTGCGTACGCCCAGT
CCCCACGACAGTCCGGTTTGTAGATTCCCTGATCTGCAATTCTTCCCGTTCCTTCATGGATTGAAGGCTCTCCT
TTCTTCCTTGAATGACTTTGCATCCCTCTCGTTTGCTGAGAGTTGGGACAAATGTTGGATTACTGGTGGAACCAAG
CCCACCACATACTGTAAATACACTCTTCTGACCAATGACCTGACTGAGGAAGTGATGGAGGAGGTGCTGCAAAA
GAAGGCAGACCTCATTCTCTCTACCATCCGCCTATCTTCCGACCCATGAAGCGCATAACCTGGAACACATGGAA
GGAGCGCCTGGTGATCCGGGCTCTGGAGAACAGAGTCGGTATCTACTCTCCTCATAACAGCCTATGATGCTGCGCC
CCAGGGCGTCAACAACCTGGTTGGCTAAAGGGCTTGGAGCTTGACCTCCAGGCCCATAACATCCTTCCAAAGCTCC
CAACTACCTACAGAGGGAAACCACCGAGTAGAATTCAACGTTAACTACACCCAAGACCTGGACAAAGTCATGTC
TGCAGTGAAAGGAATTGACGGTGTCTTCTGTCACTTCTTTTTCTGCTAGGACTGGTAATGAGGAACAAACACGGAT
TAATCTGAATTGTACTCAGAAGGCTTTGATGCAGGTGGTAGATTTTCTTTCCCGGAACAAACAACCTTTATCAGAA
GACGGAAATTCTGTCACTGGAGAAGCCTTTGCTTCTACATACTGGAATGGGACGGTTATGCACACTGGATGAATC
TGTCTCCCTGGCAACCATGATTGATCGAATAAAAAGACACCTAAAACTATCTCATATTTCGCTTAGCCCTTGGGGT
GGGGAGAACCTTAGAGTCTCAAGTCAAAGTCGTGGCCCTGTGTGCTGGTTCTGGGAGCAGCGTTCTGCAGGGTGT
TGAGGCTGACCTTTACCTCACAGGTGAGATGTCCCATCATGATACTTTGGATGCTGCTTCCCAAGGAATAAATGT
CATCCTCTGTGAACACAGCAACACTGAACGAGGCTTTCTTTCTGACCTTCGAGATATGCTGGATTCTCACTTGA
GAATAAGATAAATATTATCCTATCAGAGACTGACAGGGACCCCTTTCAGGTGGTATAATTGCAGAAACATCAGGA
TAACACATTCTACAAATCAGCTGGATGCCAACTTAAATTTGTAAACATGAGTCAGTGGGACTGGTGTGCTTCCAGA
GAGTGTCTTCGAGGGTATCATCATTTCCGGTTTGTAAATCTTATTCACCAAATGTTCTATCGCTCGTAAGGTAAA
ACTGTAATATAACTACCATATTAAATAACAAATGTTTCATTATAAACTCTAGGAAAGATTGAATAAAATCTGTTTA
CTTAACATTC

1004/6881
FIGURE 937

MLSSCVRPVPTTVRFVDSLICNSSRSFMDLKALLSSLNDFASLSFAESWDNVGLLVEPSPPHTVNTLFLTNDLTE
EVMEEVLQKKADLILSYHPPIFRPMKRITWNTWKERLVIRALENRVGIYSPHTAYDAAPQGVNNWLAKGLGACTS
RPIHPSKAPNYPTTEGNHRVEFNVNYTQDLQVMSAVKGIDGVSVTSFSARTGNEEQTRINLNCTQKALMQVVDL
SRNKQLYQKTEILSLEKPLLLHTGMGRCLTLDSEVSLATMIDRIKRHLKLSHIRLALGVGRTLESQVKVVALCAG
SGSSVLQGVEADLYLTGEMSHHDTLDAASQGINVILCEHSNTERGFLSDLRDMLDSHLENKINIILSETDRDPLQ
VV

1005/6881
FIGURE 938

CTCCTTTCTGCCCCTGGACGCCGCCGAAGAAGCATCGTTAAAGTCTCTCTTCACCCCTGCCGTCATGCTCTAAGTCA
GAGTCTCCTAAAGAGCCCCGAACAGCTGAGGAAGCTCTTCATTGGAGGGTTGAGCTTTGAAACAACCGATGAGAGC
CTGAGGAGCCATTTTGTAGCAACGGGGAACGCTCACAGACTGTGTGGTAATGAGAGATCCAAACACCAAGTGCTCC
ACGGGCTTTGGGTTTGTACATATGCCACTGTGAAGGAGGTGGAGGCAGCTATGAATGCAAGGCCACAGAAGGTG
GATGGAAGAGTCGTGGAACCAAGAGAGCTGTCTCGAGAGAAGATTCTCAAAGACCAGGTGCCACTTAACTGTG
AAAAAGATATTTGTTGGTGGCATTAAAGAAGACACTGAAGAACATCACCTAAGAGATTATTTTGAACAGTATGGA
AAAATTGAAGTGATTGAAATCATGACTGACCGAGGCAGTGGGAAGAAAAGGGGCTTTGCCTTTGTAACCTTTGAT
GACCATGACTCCGTGGATAAGATTGTCAATCAGAAATACCACACTGTGAATGGCCACAACCTGTGAAGTTAGAAAA
GCCCTGTCAAAGCAAGAGATGACTAGTGCTTCATCTAGCCAAAGAGGTGGAAGTGGTTCTGGAACTTTGGTGGT
GGTCGTGGAGGTGGTTTTAGTGGGAATGACAACCTTTGGTCATGGAAGAACTTCAGTGGTCATGGTGGCTTTGGT
GGCAGCCGTGGTGGTGGTGGATATGATGGCAGTGGGGATGGCTATAATGGATTGGTAATGATGGAAGCCATTTT
GGAGGTGGTGAAGCTACAATGATTTTGGCAATTACAAAAATCAGTCTTCAAATTTTGGACCCGTGAAGGGAGGA
AATTTTGGAGGCAGAAGCTCTGGCCCCATATGGCGGTGGAGGCCAATACTTTGCAAACCACGAACTAAGGTGGC
TATGGCGGTTCCAGCAGTAGCAGTAGCTATGGCAGTGGCAGAAGATTTTAAATTAGGAAACAAAGCTTAGCAGGAG
AGGAGAGCCAGAGAAGTGACAGGGGAAGCTACAGGTTACAACAGATTTGTGAACTCAGCCAAGCACAGTGGTGGCA
GGGCCTAGCTGCTACAAAGAAGACATGTTTTAGACAAATACTCATGTGTATGGGCAAAAACTCGAGGACTGTAT
TTGTGACTAATTGTATAACAGGTTATTTTAGTTTCTGTTCTGTGGAAAGTGTAAGCATTCCAACAAAGGGTTTT
AATGTAGATTTTTTTTTTGCACCCATGCTGTTGATTGCTAAATGTAATAGTCTGATCGTGACGCTGAATAAATG
TCTTTTTTTTTTAAATGTGCTGTGTAAAGTTAGTCTACTCTGAAGCCATCTTGGTAAATTTCCCCAACAGTGTGAAG
TTAGAATTCCTTCAGGGTGATGCCAGGTTCTATTTGGAATTTATATACAACCTGCTTGGGTGGAGAAGCTATTGT
CTTCGGAACCTTGGTGTAGTTGAACTGATAGTTACTGTTGTGACCTGAAGTTCACCATTAAAAGGGATTACCCA
AGCAAAATCATGGAATTATTGGTTATAAAAGTGATTGTTGGCACATCCTATGCAATATAICTAAATTGAATAATG
GTACCAGATAAAATTATAGATGGGAATGAAGCTTGTGTATCATCCATTATCATGTGTAATCAATAACGATTTAA
TTCTCTGGAA

1006/6881
FIGURE 939

MSKSESPKEPEQLRKLF IGGLSFETTDESLRSHFEQRTLTDCVVMRDPNTKCSTGFGFV TYATVKEVEAAMNAR
PQKVDGRVVEPKRAVSREDSQRPGAHLTVKKIFVGGIKEDTEEHHLRDYFEQYKIEVIEIMTDRSGGKKRGFAF
VTFDDHDSVDKIVIQKYHTVNGHNCEVRKALSKQEMTSASSSQRGRSGGNFGGGRGGGFSGNDNFGHGRNFSGH
GGFGGSRGGGGYDGS GDGYNGFGNDGSHFGGGGSYNDFGNYKNQSSNFGPVKGGNFGGRSSGPYGGGGQYFAKPR
NRWLWRFQQVAVAMAVAEDF

1007/6881
FIGURE 940

GAGGCGTAGCGGAAGTTACTGCAGCCGCGGTGTTGTGCTGTGGGAAGGGAGAAGGATTTGTAAACCCCGGAGCG
AGGTTCTGCTTACCCGAGGCCGCTGCTGTGCGGAGACCCCGGGTGAAGCCACCGTCATCATGTCTGACCAGGAG
GCAAACCTTCAACTGAGGACTTGGGGGATAAGAAGGAAGGTGAATATATTAACTCAAAGTCATTGGACAGGAT
AGCAGTGAGATTCACCTCAAAGTGAAATGACAACACATCTCAAGAACTCAAAGAATCATACTGTCAAAGACAG
GGTGTTCGAATGAATTCACCTCAGGTTTCTCTTTGAGGGTCAGAGAATTGCTGATAATCATACTCCAAAAGAACTG
GGAATGGAGGAAGAAGATGTGATTGAAGTTTATCAGGAACAAACGGGGGGTCATTCAACAGTTTAGATATTCTTT
TTATTTTTTTCTTTCCCTCAATCCTTTTTTATTTTTTAAAAATAGTTCTTTTGTAATGTGGTGTTCAAAACGGA
ATTGTTATTCATTATTGTTTGTTCATTGTGCTGATTTTGGTGATCAAGCCTCAGTCCCCTTCATATTACC

1008/6881
FIGURE 941

ATGTTGGTGCTGTTTTGAAACGCTCTGTGGGTACGCCATCTTTAAGGTTCTAAATGAGAAGAACTTCAAGAGGTT
GATAGTTTATGGAAAGAATTTGAAACTCCAGAGAAAGCAAACAAAATAGTAAAGCTAAAACATTTTGAGAAATTT
CAGGATACAGCAGAAGCATTAGCAGCATTACAGCTCTGATGGAGGGCAAAATCAATAAGCAGCTGAAAAAGTT
CTGAAGAAAATAGTAAAAGAAGCCCATGAACCGCTGGCAGTAGCTGATGCTAAACTAGGAGGGGTCATAAAGGAA
AAGCTGAATCTCAGTTGTATCCATAGTCCTGTTGTTAATGAACCTTATGAGAGGAATTCGTTTACAAATGGATGGA
TTAATCCCTGGGGTAGAACCACGTGAAATGGCAGCTATGTGTCTTGGATTGGCTCACAGCCTGTCTCGATATAGA
TTGAAGTTTAGCGCTGATAAAGTAGACACAATGATTGTTTCAGGCAATTTCCCTGTTAGATGACTTGGATAAAGAA
CTAAACAACACTACATTATGCGATGTAGAGAATGGTATGGCTGGCATTTCCTGAATTAGGAAAAATTATTTTCAGAT
AATTTAACATACTGCAAGTGTTTACAGAAAGTTGGCGATAGGAAGAACTATGCCTCTGCCAAGCTTTCTGAGTTG
CTGCCAGAAGAAGTTGAAGCAGAAGTGAAAGCAGCTGCAGAGATATCAATGGGAACAGAGGTTTCAGAAGAAGAT
ATTTGCAATATTCTGCATCTTTGCACCCAGGTGATTGAAATCTCTGAATATCGAACCCAGCTCTATGAATATCTA
CAAAATCGAATGATGGCCATTGCACCCAATGTTACAGTCATGGTTGGGGAATTAGTTGGAGCACGGCTTATTGCT
CATGCAGGTTCTCTTTTAAATTTGGCCAAGCATGCAGCTTCTACCGTTTCAGATTCTTGGAGCTGAAAAGGCACTT
TTCAGAGCCCTCAAATCTAGACGGGATACCCCTAAGTATGGTCTCATTATCATGCTTCACTCGTGGGCCAGACA
AGTCCCAAACACAAAGGAAAGATTTCTCGAATGCTGGCAGCCAAAACCGTTTTGGCTATCCGTTATGATGCTTTT
GGTGAGGATTCAAGTTCTGCAATGGGAGTTGAGAACAGAGCCAAATTAGAGGCCAGGTTGAGAACTTTGGAAGAC
AGAGGGATAAGAAAAATAAGTGGAACAGGAAAAGCATTAGCAAAAACAGAAAAATATGAACACAAAAGTGAAGTG
AAGACTTACGATCCTTCTGGTGACTCCACACTTCCAACCTGCTCTAAAAAACGCAAAATAGAACAGGTAGATAAA
GAGGATGAAATTACTGAAAAGAAAGCCAAAAAGCCAAGATTAAAGTTAAAGTTGAAGAAGAGGAAGAAGAAAAA
GTGGCAGAAGAAGAAGAAACATCTGTGAAGAAGAAGAAGAAAGGGGTAAAAAGAAACACATTAAGGAAGAACCA
CTTTCTGAGGAAGAACCATGTACCAGCACAGCAATTGCTAGTCCAGAGAAAAAGAAGAAAAAGAAAAAAGAGA
GAGAACGAGGATTAA

1009/6881
FIGURE 942

GGAGGCCAAGGTGCAACTTTCTTCGGTCGTCCCGAATCCGGGTTTCATCCAACACCAGCCGCCTCCACCATGCCGC
CGAAGTTCGACGGTGCCACTTCTGCCCTGGCCCCCAAGATCGGCCCCCTGGGTCTGTCTCCAAAAAGGTTGGTG
ATGACATTGCCAAGGCAACGGGTGACTGGAAGGGCCTGAGGATTACAGTGAAACTGACCATTGAGAACAGACAGG
CCCAGATTGAGGTGGTGCCTTCTGCCTCTGCCCTGATCATCAAAGCCCTCAAGGAACCAAGAGACAGAAAGA
AACAGAAAAACATTAAACACAGTGGGAATATCACTTTTGATGAGATCGTCAACATTGTTGACAGATGCGGCACC
GATCCTTAGCCAGAGAACTCTCTGGAACCATTAAAGAGATCCTGGGGACTGCCCAGTCTGTGGGCTGTAATGTTG
ATGGCCGCCATCCTCATGACATCATCGATGACATCAACAGTGGTGCTGTGGAATGCCCAGCCAGTTAAGCACAAA
GGAAATATTTCAATAAAGGATCATTTGACAAC TG

1010/6881
FIGURE 943

MPPKFDGATSALAPKIGPLGLSPKKVGDDIAKATGDWKGLRITVKLTIQNRQAQIEVVPSASALI IKALKEPPRD
RKKQKNIKHSGNITFDEIVNIVRQMRHRSLARELSGTIKEILGTAQSVGCNVDGRHPHDIIDDINS GAVECPAS

1011/6881
FIGURE 944

ATGGAAGATTGATGGACATGGACATGAGCCTCCTGAGGCCCCAGAACTATCTTTTCGGTTGTGAACTAAAGGCC
GACAAAGATGATCACTTTAAGGTGGATAATGATGAAAATGAGACCATCAGTTTAGGGGCTGGTGCAAGGGATGAA
TTGCACATTGTTGAAGCAGAGGCCATGAATGATGAAGGCAGTCCAATTAAAGTCACACTGGCAACTTTGAAAATG
TCTGTACAGCCAACCTGCTTCCCTTGGGGGCTTTGAAATCACACCACCAGTGGTCTTACAGTTGAAGTGTGGTTCA
GGGCCAGTGCATATTAGTAGACAGTGCTTAGTAGCTGTGGAGGAAGATGCAGAGTCAGAAGATGAAGAGGAGGAG
GATGTGAACCTCTTAAGCATATCTGGAAGCGGTCTGCCTCTGGAGGTGGTAGCAAGGCGCCAGCTACCGCGCC
CACTGGTCAGAAGTGTGGCCAGCGTCTATGCAGGCCCTGGGAGCTCAGGGTGGCTTGGGGTCCAGGAGCATGGCC
GCAGGTATGGCCGGGGTCTGGCAGGAATGGGAGGCATCCAGAACGAGAAGGAGACCATGCAAAGCCTGAATGAC
CGCCTGGCCTCCTACCTGGACAGAGTGAGGAGCCTGGAGATGGGAACTGGAAGCTGGAGAGCAAAATCTGGGAG
CACCTGGAGAAGAAGGGACCCAGGTGAGAGCTGGGGCCACTACTTCAAGACCATCGAGGAGGACCTGACTCAG
ATCTTCACAAGTACTGTGGACAATACCTGCATCATTCTGCAGATCGACAATGCCATCTTGCTGCTGATGACTTT
AGAGTCAAGTATGAGACAGAGCCGGCCACGTGCCAGTCTGTGGAGAACGACATCCATGGGCTCCACAAGGTCATT
GATGACACCAGTGTCACTCAGCTGCAGCTAGAGACAGAGATCGAGGCTCTCAAGGAGGAGCTGCTCTTCCTGAAG
AAGAACCACGAAGAGGAAGTAAAAGGCCTACAAGCCCAGATTGCCAGCTCTGTTGACCATGGAGGCCTTGAGAT
CGACCTGGACTGGATGAAAAATCTGGAGGCCAGCTTGGAGAACAGCCTGAAGGGGGGGTGGAGGCCTGCTACACC
CTGCAGATGGAACAACCTCAATGGGATCCTGCTGCACCTGGAGTCAGAGCTGGCACAGACCTGGGCAGAGGGACAG
GGCCAGGCCCCAGGAGTACCAGGCCCTGCCAAACATCAAGCAGCAACTCCACGCAAACCATCCAAAAGACTCCAC
CTGCCAACAGTGGATGGCAAAGTGGTGTCTGAGACCAATGACACCAAAGTTCTGAGACATTAA

1012/6881
FIGURE 945

MEDLMDMDMSLLRPQNYLFGCELKADKDDHFKVDNDENETISLGAGARDELHIVEAEAMNDEGSPIKVTLATLKM
SVQPTASLGGFEITPPVVLQLKCGSGPVHISRQCLVAVEEDAEESEDEEEEDVNLLSISGKRSASGGGSKAPSYRA
HWSEVWPASMQALGAQGGLGSRSMAGMAGGLAGMGGIQNEKETMQSLNDRLASYLDRVRSLEMGNWKLESKIWE
HLEKKGPQVRDWGHYFKTIEEDLTQIFTSTVDNTCIIILQIDNAHLAADD FRVKYETEPATCQSVENDIHGLHKVI
DDTSVTQLQLETEIEALKEELLFLKKNHEEEVKGLQAQIASSVDHGGLGDRPGLDEKSGGQLGEQPEGGVEACYT
LQMEQLNGILLHLESELAQTWAEQGQQAQEQALPNIKQQLHANHPKDSHLPTVDGKVVSETNDTKVLRH

1013/6881
FIGURE 946

GACAGTTT TAGCAGAACAGCCTCCGCGGCTCCGGGGAGAAGCAATATGTTAAGGATACCTGTAAGAAAGGCCTTAG
TAGGCCTTTCTAAGTCTCTCTAAAGGATGTGTTCTGAACAACCTGCCACAGCAGCAAGCAACTTGATTGAAGTATTTG
TTGATGGTCAGTCTGTCTATGGTGGAAACCGGGAACGACCGTCTCCAGCTTGTGAGAAGGTTGGCATGCAGATCC
CTCGATTCTGTTATCATGAAAGGTTGTCTGTTGCTGGAAACTGCAGGATGTGCCTTGTTGAAATTGAGAAAGCCC
CTAAGGTTGTAGCTGCTTGTGCCATGCCAGTAATGAAGGGTTGGAATATCCTAACAACTCAGAAAAATCCAAAA
AAGCCAGGGAAGGTGTGATGGAGTTCTTATTAGCAATCACCCATTGGACTGTCCTATTTGTGACCAGGGAGGTG
AATGTGATCTGCAGGACCAGTCCATGATGTTTGGAAATGATAGGAGCCGATTTTTAGAGGGGAAGCGTGCTGTGG
AAGACAAGAACATTGGGCCATTGGTAAAGACCATCATGACAAGATGTATACAGTGTACTCGCTGCATCAGGTTTG
CAAGTGAGATTGCAGGAGTAGATGATTGGGAACAACAGGCAGAGGAAATGATATGCAAGTTGGCACATACATTG
AAAAGATGTTTCATGTCTGAACTGTCTGGGAATATCATTGATATCTGCCCTGTAGGTGCCCTAACCTCTAAGCCCT
ATGCCTTTACTGCCCGGCTTGGGAAACAAGAAAGACAGAATCCATTGATGTAATGGATGCGGTTGGAAGTAATA
TTGTGGTTAGCACAAGAACTGGAGAAGTGATGAGGATTTGCCACGTATGCATGAGGACATCAATGAAGAGTGGA
TCTCTGATAAAACCAGATTTGCCTATGATGGGCTAAAACGTCAAAGACTTACCGAGCCAATGGTCAGAAATGAAA
AAGGGCTTTTAACTATACTTCTTGGGAGGATGCGCTCTCTCGCGTAGCTGGAATGTTGCAGAGTTTTCAAGGCA
AAGATGTGGCAGCAATTGCAGGTGGCTTGGTGGATGCTGAAGCCCTGGTAGCTCTCAAAGATTTGCTTAATAGAG
TGGACTCTGACACCTTATGCACTGAAGAGGTCTTCCCCACTGCAGGAGCTGGCACAGATTTGCGTTCCAATTATC
TTCTTAATACTACAATTGCTGGTGTGGAAGAGGCAGATGTTGTTCTTCTGTTGGTACAAACCCACGTTTTGAGG
CACCCTGTTTAAATGCTAGAATTCGAAAGAGCTGGCTGCATAATGACTTAAAAGTGGCCCTTATAGGCAGTCCAG
TGGACCTCACTTACACATATGACCACCTGGGAGACTCCCCAAAATTCTTCAAGACATTGCTTCGGGAAGCCATC
CATTAGCCAGGTCTTAAAGGAAGCTAAAAAACCAATGGTGGTTTTAGGCAGTTCTGCACTCCAAAGAAATGATG
GAGCAGCAATTCTTGCACTGTTTTCTAGCATTGCACAAAAGATTCCGGATGACTAGTGGTGTACTGGTGATTGGA
AAGTTATGAATATCCTTCATAGGATTGCAAGTCAAGTAGCTGCTTTGGACCTTGGCTATAAGCCTGGGGTGGAG
CAATTCGGAAGAACCCTCCCAAGGTGCTGTTTTCTCTGGGAGCAGATGGAGGTTGTATCACACGACAGGATTTGC
CAAAGGATTGTTTCATTATTTATCAAGGACATCATGGTGATGTTGGGGCTCCCATAGCTGATGTTATTCTCCAG
GAGCTGCTTACACAGAGAAGTCTGCTACATATGTCAACACTGAGGGTAGAGCTCAGCAGACTAAGGTAGCAGTGA
CACCTCCTGGCTTGGCAAGAGAAGACTGGAAAATTATAAGAGCACTCTCTGAGATTGCTGGAATGACTCTTCCAT
ATGATACTCTGGATCAAGTAAGGAACAGATTGGAAGAAGTCTCTCCTAATCTTGTTGATATGATGATATTGAAG
GGGCTAATTACTTCCAGCAAGCAAATGAGCTCTCAAAGCTAGTGAACCAGCAGCTTCTTGCTGACCCACTTGTTT
CACCTCAGCTAACTATAAAAGACTTCTACATGACAGATTCAATTAGCAGAGCCTCACAGACAATGGCCAAATGTG
TCAAAGCTGTACAGAGGGTGCCAGGCAGTAGAGGAACCATCCATATGCTGAAGCTTCTACTAGGATCCCAGTT
TTGCCGCGAGATAATTAATGGACAACGTAGTGCAGTGATCCTTTACAGGTTTATTTCTTTGTAAAAAAAATAAT
AATAATTTGAATCATGTAATATTTAAGGTTATACTATGCCTATTTGAAAATGATATTAGTTATCACTTTGCAGT
TTGAAAAACATGTATTGTGTAAAGGTTAAATAACAAACTATGCAGATGCTCTTAAAGCATTGATAACCTTT
GTGACGAACATAAAGAGATCCTTAAATT

1014/6881
FIGURE 947

MLRIPVRKALVGLSKSPKGCVRRTATAASNLIIEVFVDGQSVMEPGTTVLQACEKVGMQIPRFCYHERLSVAGNC
RMCLVEIEKAPKVVAACAMPVMKGWNILTNSEKSKKAREGVMEFLLANHPLDCPICDQGGECDLQDQSMFNGNR
SRFLEGKRAVEDKNIGPLVKTIMTRCIQCTRCIRFASEIAGVDDLGTTRGRGNDMQVGTYYIEKMFMSSELGNIIDI
CPVGALTSKPYAFTARPWETRKTESIDVMDAVGSNIVVSTRTGEVMRILPRMHEDINEEWISDKTRFAYDGLKRQ
RLTEPMVRNEKGLLTYTSWEDALSRVAGMLQSFQGKDVAIAGGLVDAEALVALKDLLNRVDSDTLCTEEVFPTA
GAGTDLRSNYLLNTTIAGVEEADVLLVGTNPRFEAPLFNARIRKSWLHNDLKVALIGSPVDLTYTYDHLGDSPK
ILQDIASGSHPFSSQVLKEAKKPMVVLGSSALQRNDGAAILAAVSSIAQKIRMTSGVTGDWKVMNILHRIASQVAA
LDLGYKPGVEAIRKNPPKVLFLLGADGGCITRQDLPKDCFIYQGHGHDVGAPIADVILPGAAYTEKSATYVNT
GRAQOTKVAVTPPGLAREDWKIIRALSEIAGMTLPYDTLDQVRNRLEEVSPNLVRYDDIEGANYFQQANELSKLV
NQQLLADPLVPPQLTIKDFYMTDSISRASQTMACVKAVTEGAQAVEEPSIC

1015/6881
FIGURE 948

GCCTCGAGGCGGGCGTCTTCGGTCATCTCCGGCGCTTCTAGGGCTGGTTCCCGTCATCTTCGGGAGCCGTGGAGC
TCTCGGATACAGCCGACACCATGGGTTTCGGAGACCTGAAAAGCCCTGCCGGCCTCCAGGTGCTCAACGATTACC
TGGCGGACAAGAGCTACATCGAGGGGTATGTGCCATCACAAGCAGATGTGGCAGTATTTGAAGCCGTGTCCAGCC
CACCGCCTGCCGACTTGTGTCATGCCCTACGTTGGTATAATCACATCAAGTCTTACGAAAAGGAAAAGGCCAGCC
TGCCAGGAGTGAAGAAAGCTTTGGGCAAATATGGTCCTGCCGATGTGGAAGACACTACAGGAAGTGGAGCTACAG
ATAGTAAAGATGATGATGACATTGACCTCTTTGGATCTGATGATGAGGAGGAAAGTGAAGAAGCAAAGAGGCTAA
GGGAAGAACGTCCTTGACAAATATGAATCAAAGAAAGCCAAAAACCTGCACTTGTTGCCAAGTCTTCCATCTTAC
TAGATGTGAAACCTTGGGATGATGAGACAGATATGGCGAAATTAGAGGAGTGCCTCAGAAGCATTCAAGCAGACG
GCTTAGTCTGGGGCTCATCTAACTAGTTCCAGTGGGATACGGAATTAAGAACTTCAAATACAGTGTGTAGTTG
AAGATGATAAAGTTGGAACAGATATGCTGGAGGAGCAGATCACTGCTTTTGAGGACTATGTGCAGTCCATGGATG
TGGCTGCTTTCAACAAGATCTAAAATCCATCCTGGATCATGGCATTTAAATAAAAGATTGAAAGATTAAAA

1016/6881
FIGURE 949

MGFGDLKSPAGLQVLNDYLADKSYIEGYVPSQADVAVFEAVSSPPPADLCHALRWYNHIKSYEKEKASLPGVKKA
LGKYGPADVEDTTGSGATDSKDDDDIDLFGSDDEEESEEAKRLREERLAQYESKKAKKPALVAKSSILLDVKPWD
DETDMAKLEECVRSIQADGLVWGSSKLVFVGYGIIKLQICVVEDDKVGTDMLEEQITAFEDYVQSMDVAAFNKI

1017/6881
FIGURE 950

GGACAATTTGTGGGCCATTTAATTCAGGGCCCCCAATTCGTACGTGGAGAAGTGGGAATGCAAAAGTACTTTGAC
CTTTAACCTTCGGTCCGGCGCGGTGGAGGGAAACGCCTCCGTCTCTATATAAGGAATTTTCCGGTCTCTTCGGGT
CCTTTTTCCTCTCTTCAGCGTGGGGCGCCCAACAATTTGCGCGCTCTCTTTCTGCTGCTCCCCAGCTCTCGGATAC
AGCCGACACCATGGGTTTCGGAGACCTGAAAAGCCCTGCCGGCCTCCAGGTGCTCAACGATTACCTGGCGGACAA
GAGCTACATCGAGGGGTATGTGCCATCACAAGCAGATGTGGCAGTATTTGAAGCCGTGTCCAGCCCACCGCCTGC
CGACTTGTGTCATGCCCTACGTTGGTATAATCACATCAAGTCTTACGAAAAGGAAAAGGCCAGCCTGCCAGGAGT
GAAGAAAGCTTTGGGCAAATATGGTCCTGCCGATGTGGAAGACACTACAGGAAGTGGAGCTACAGATAGTAAAGA
TGATGATGACATTGACCTCTTTGGATCTGATGATGAGGAGGAAAGTGAAGAAGCAAAGAGGCTAAGGGAAGAACG
TCTTGACAATATGAATCAAAGAAAGCCAAAAAACCTGCACCTTGTTGCCAAGTCTTCCATCTTACTAGATGTGAA
ACCTTGGGATGATGAGACAGATATGGCGAAATTAGAGGAGTGCCTCAGAAGCATTCAAGCAGACGGCTTAGTCTG
GGGCTCATCTAACTAGTTCCAGTGGGATACGGAATTAAGAACTTCAAATACAGTGTGTAGTTGAAGATGATAA
AGTTGGAACAGATATGCTGGAGGAGCAGATCACTGCTTTTGAGGACTATGTGCAGTCCATGGATGTGGCTGCTTT
CAACAAGATCTTAAAATCCATCCTGGATCATGGCATTATAAATAAAAGATTGAAAGATTAAAA

1018/6881
FIGURE 951

MGFGDLKSPAGLQVLNDYLADKSYIEGYVPSQADVAVFEAVSSPPPADLCHALRWYNHIKSYEKEKASLPGVKKA
LGKYGPADVEDTTGSGATDSKDDDDIDLFGSDDEEESEEAERLREERLAQYESKKAKKPALVAKSSILLDVKPWD
DETDMAKLEECVRSIQADGLVWGSSKLVPVGYGIKKLQICVVEDDKVGTDMLEEQITAFEDYVQSM DVAAFNKI

1019/6881
FIGURE 952

GAAGGGGCGGGGCCAAAACCTGCGCGCCCAATCGGGGTGACGCTCTAGCCTTGCCGGGGACTCGTGGGTAACTTGC
TTTTGGGAGCCAGCGGTATGGCGTCGGGCTGCAAGATTGGCCCGTCCATCCTCAACAGCGACCTGGCCAATTTAG
GGGCGGAGTGCTCCGGATGCTAGACTCTGGGGCCGATTATCTGCACCTGGACGTAATGGACGGGCATTTTGTTC
CAACATCACCTTTGGTCACCTGTGGTAGAAAGCCTTCGAAAGCAGCTAGGCCAGGACCCTTTCTTTGACATGCA
CATGATGGTGTCCAAGCCAGAACAGTGGGTAAAGCCAATGGCTGTAGCAGGAGCCAATCAGTACACCTTTTCATCT
CGAGGCTACTGAGAACCCAGGGGCTTTGATTAAAGACATTCCGGGAGAATGGGATGAAGGTTGGCCTTGCCATCAA
ACCAGGAACCTCAGTTGAGTATTTGGCACCATGGGCTAATCAGATAGATATGGCCTTGGTTATGACAGTGGAAACC
GGGGTTTGGAGGGCAGAAATTCATGGAAGATATGATGCCAAAGGTTCACTGGTTGAGGACCCAGTTCCCATCTTT
GGATATAGAGGTGCGATGGTGGAGTAGGTCCTGACACTGTCCATAAATGTGCAGAGGCAGGAGCTAACATGATTGT
GTCTGGCAGTGCTATTATGAGGAGTGAAGACCCAGATCTGTGATCAATCTATTAAGAAATGTTTGCTCAGAAGC
TGCTCAGAAACGTTCTCTTGATCGGTGAACCAATAAGGAGCCAGTGTTTCTGTTTCATGAAATCTCCCTTTTACT
GGAAACAGGAATATTGACTACCAATCACAATGCAATTGAAGCCGTAAGTCTTTTGGAGCAGTTATTTCATTCC
AGTGATTAAACTGATTGTGAGAATATTCTAAGAGGTGAGAAATGGTGTGTATAACTACATTTTATGATGTC
AATTTATTGATTAGTGAGTAAGATACTGTTTTATTGAGAGATTTGATTTTTATAAAGTAAAAATACGGCTGCAT
TAGGGTTACAAACAGAAAAGTGCTTAATGTCTAAGGAGGGCATATTAGCTACACTACAAAAACAAATTTGTCT
GTACTTCTGAAAAGAATTTTGTGTTTCTCAGCTGTTTTCCAAAAGCAAAGGAAGTCTTATGGTTTTTTCTAT
TTCATGTTATTGTGATTTGTTTATAAGTTTGGGTGGGGTGATACCATATTCTTGGTTCTTAAATCTATCACTT
TTCACCTTACACTTGATGTGTGAAAACATAAAAAACATGTGTGAAACCCAGGGGTTCTAAATACAAGCATAGA
TTTTATCAGGGTGTTTTGTCAAAGCAGGTTATTCACTGATTCCTCCCCACCATTCTTAAAGACGTTAAATAATGC
TGTTGTGTTAGCTCTGAGTAGAAAGGAAAAAGTAAACCTCTGTTTGGAGGTAATATTGGGTGAATTCTGACTG
CCCCTTTCTAGCTGGACCTTTAACAAATCACCAATCTTTTTTGTGTTTCTCTAAAGTCATTTATACATTAAATG
TAATTATAGCAACTGTGGGGTTCTGTTGAGAATTAAGAGCTAACACTATATATGTAAAGTTTCCAGTACTAGTCC
CAGAATTTAGAAATATGCTCAACACAAAAGTAAACAGCATTATATAAGTTTATATTTTGTGAGTTATAAAGTACTT
TGATATATTCTCATTAAATCTGTAAATCACCTCTATAAGTAAGTGGTAATAATAAAGCAGATATTTTTGTCCCA
TTAAAAAATGAAGAAATTAATGCTTAATAGGGTGGTACCCTGGAAAGGATCTGGGAAGTGGTAGAATTTCTGGT
CTGTACTTTTACAAATGGAGCCCTTGGGAGGTGGGTTAGGTAAAGAAGCTTTTACTTAACGTTGTCTTATTTT
CAGTCTAATTTTACGCTGTAGCAGAACCAGATGGCTGAGAAAATCTGGAACTATGGATCTTGACCCCAAGGATA
TATTATTTTATTCCAAGAAAGATCAGGTAGGCGAAAAGATGACAGGATACAGAGTCAATCCATAAACTAAATAT
TATAACTGTTCTGAATTATACAGAGTCTAAAAATATGTGTCAGCTACTTCATTCTGTAAATACTCTTGCTGTGT
TATAAATATGGCAAGAAATAAACATGACCAATATCAATAGACTTCTTGAGGCTACTATAAGTTTGGAGAAATAAG
GTTCAAAAAATAAGAAATGCTAACACTTAAGCACAGACTAGAGCTTGCTTGGGTTTCTTCTGCATTACAAGGTAA
AAATTTGTTAATGTTTGTGTTTTATTTCAGCTTGGGAAAGCTTTGTGCCATGAATACGTCGCATTAAATAACAAGCA
ACACACGGCATATAGAAATAACTTTAATAAAAAACTTACATAGAAGATTATAATATCAGACGTGACAAAGATT
GAGTTTATTTGCCTGGACAACCTTGGGTTTGTCTGGCTTTTGTGTTTCTTTTCTTTTAAAAATAAATGTACAGTAA
ACTAC

1020/6881
FIGURE 953

MLDSGADYLLHLDVMDGHFVPNITFGHPVVESLRKQLGQDPFFDMHMMVSKPEQWVKPMAVAGANQYTFHLEATEN
PGALIKDIRENGMKVGLAIKPGTSVEYLPWANQIDMALVMTVEPGFGGQKFMEDMMPKVHWLRTQFPSLDIEVD
GGVGPDTVHKCAEAGANMIVSGSAIMRSEDPRSVINLLRNVCSAAQKRSLDR

1021/6881
FIGURE 954A

GAAGAGCAAGAGGCAGGCTCAGCAAATGGTTTCAGCCCCAGTCCCCGGTGGCTGTCAGTCAAAGCAAGCCCCGGTTG
TTATGACAATGGAACACTATCAGATAAATCAACAGTGGGAGCGGACCTACCTAGGCAATGCGTTGGTTTGTAC
TTGTTATGGAGGAAGCCGAGGTTTTAACTGCGAGAGTAAACCTGAAGCTGAAGAGACTTGCTTTGACAAGTACAC
TGGGAACACTTACCGAGTGGGTGACACTTATGAGCGTCCCTAAAGACTCCATGATCTGGGACTGTACCTGCATCGG
GGCTGGGCGAGGGAGAATAAGCTGTACCATCGCAAACCGTGCCATGAAGGGGGTCAGTCTACAAGATTGGTGGA
CACCTGGAGGAGACCACATGAGACTGGTGGTTACATGTTAGAGTGTGTGTGCTTGGTAATGGAAGGAGAAATG
GACCTGCAAGCCCATAGCTGAGAAGTGTGTTGATCATGCTGCTGGGACTTCCTATGTGGTCGGAGAAACGTGGGA
GAAGCCCTACCAAGGCTGGATGATGGTAGATTGTAATTGCGCTGGGAGAAGGCAGCGGACGCATCACTTGCACCTC
TAGAAATAGATGCAACGATCAGGACACAAGGACATCCTATAGAATTGGAGACACCTGGAGCAAGAAGGATAATCG
AGGAAACCTGCTCCAGTGCATCTGCACAGGCAACGGCCGAGGAGAGTGGAAAGTGTGAGAGGCACACCTCTGTGCA
GACCACATCGAGCGGATCTGGCCCCCTTACCAGTGTTCGTGCAGCTGTTTACCAACCGCAGCCTCACCCTCAGCC
TCCTCCCTATGGCCACTGTGTACAGACAGTGGTGTGGTCTACTCTGTGGGATGCAGTGGCTGAAGACACAAGG
AAATAAGCAAATGCTTTGCACGTGCCTGGGCAACGGAGTCAGCTGCCAAGAGACAGCTGTAACCCAGACTTACGG
TGGCAACTCAAATGGAGAGCCATGTGTCTTACCATTACCTACAATGGCAGGACGTTCTACTCTGCACCACAGA
AGGGCGACAGGACGGACATCTTGGTGCAGCACAACTTCGAATTATGAGCAGGACCAGAAATACTCTTCTGCAC
AGACCACACTGTTTTGGTTTCAGACTCGAGGAGGAAATTCGAATGGTGCCTTGTGCCACTTCCCCTTCTATACAA
CAACCACAATTACACTGATTGCACTTCTGAGGGCAGAAGAGACAACATGAAGTGGTGTGGGACCACACAGAATA
TGATGCCGACCAGAAAGTTTGGGTTCTGCCCCATGGCTGCCACGAGGAAATCTGCACAACCAATGAAGGGTGCAT
GTACCGCATTTGGAGATCAGTGGGATAAGCAGCATGACATGGGTACATGATGAGGTGCACGTGTGTTGGGAATGG
TCGTGGGGAATGGACATGCATTGCCTACTCGCAGCTTCGAGATCAGTGCAATTGTTGATGACATCACTTACAATGT
GAACGACACATTCCACAAGCGTCTGAAGAGGGGCACATGCTGAACGTACATGCTTCGGTCAGGGTCGGGGCAG
GTGGAAGTGTGATCCCGTCGACCAATGCCAGGATTCAGAGACTGGGACGTTTTATCAAATTGGAGATTTCATGGGA
GAAGTATGTGCATGGTGTGAGATACCAAGTGTCTACTGCTATGGCCGTGGCATTGGGGAGTGGCATTGCCAACCTTT
ACAGACCTATCCAAGCTCAAGTGGTCTGTGCAAGTATTTATCACTGAGACTCCGAGTCAGCCCCAACTCCCACCC
CATCCAGTGGAAATGCACCACAGCCATCTCACATTTCCAAGTACATTCTCAGGTGGAGACCTAAAAATTCTGTAGG
CCGTTGGAAGGAAGCTACCATACAGGCCACTTAACTCCTACACCATCAAAGGCCTGAAGCCTGGTGTGGTATA
CGAGGGCCAGCTCATCAGCATCCAGCAGTACGGCCACCAAGAAGTACTCGCTTTGACTTCACCACCACAGCAC
CAGCACACCTGTGACCAGCAACACCGTGACAGGAGAGACGACTCCCTTTTCTCCTCTTGTGGCCACTTCTGATC
TGTGACCGAAATCACAGCCAGTAGCTTTGTGGTCTCCTGGGTCTCAGCTTCGGACACCGTGTCTGGGATTCCGGGT
GGAATATGAGCTGAGTGAGGAGGGAGATGAGCCACAGTACCTGGATCTTCCAAGCACAGCCACTTCTGTGAACAT
CCCTGACCTGCTTCTGGCCGAAAATACATTGTAAATGTCTATCAGATATCTGAGGATGGGGAGCAGAGTTTGAT
CCTGTCTACTTCACAAACAACAGCGCTGATGCCCCCTCTGACACGACTGTGGACCAAGTTGATGACACCTCAAT
TGTGTTGCTGCTGGAGCAGACCCAGGCTCCCATCACAGGTACAGAATAGTCTATTCGCCATCAGTAGAAGGTAG
CAGCACAGAACTCAACCTTCTGAACTGCAAACTCCGTCAACCTCAGTGACTTGCAACCTGGTGTTCAGTATAA
CATCACTATCTATGCTGTGGAAGAAAATCAAGAAAGTACACCTGTTGTCAATCAACAAGAAACCACTGGCACCCC
ACGCTCAGATACAGTGCCCTCTCCCAGGGACCTGCAGTTTGTGGAAGTGACAGACGTGAAGGTCACCATCATGTG
GACACCGCCTGAGAGTGAGTGACCGGCTACCGGTGTGGATGTGATCCCCGTCAACCTGCCTGGCGAGCACGGGCA
GAGGCTGCCCATCAGCAGGAACACCTTTGCAGAAGTACCGGGCTGTCCCCCTGGGGTCACCTATTACTTCAAAGT
CTTTGCAGTGAGCCATGGGAGGGAGAGCAAGCCTCTGACTGCTCAACAGACAACCAAACTGGATGCTCCCACTAA
CCTCCAGTTTGTCAATGAACTGATTCTACTGTCTGCTGAGATGGACTCCACCTCGGGCCCAGATAACAGGATA
CCGACTGACCGTGGGCCCTTACCCGAAGAGGACAGCCAGGAGTACAATGTGGGTCCCTCTGTCTCCAAGTACCC
ACTGAGGAATCTGCAGCCTGCATCTGAGTACACCGTATCCCTCGTGGCCATAAAGGGCAACCAAGAGAGCCCCAA
AGCCACTGGAGTCTTTACCACACTGCAGCTGGGAGCTCTATTCCACCTTACAACACCGAGGTGACTGAGACCAC
CATTGTGATCACATGGACGCTGCTCCAAGAATTGGTTTTAAGCTGGGTGTACGACCAAGCCAGGGAGGAGAGGC
ACCACGAGAAGTGACTTCAGACTCAGGAAGCATCGTTGTGTCCGGCTTGAAGTCCAGGAGTAGAATACGTCTACAC
CATCCAAGTCTGAGAGATGGACAGGAAAGAGATGCGCCAATTGTAAACAAAGTGGTGACACCATTGTCTCCACC
AACAACTTGCATCTGGAGGCAACCCCTGACACTGGAGTGCTCAGTCTCCTGGGAGAGGAGCACCACCCAGAA
CATTACTGGTTATAGAATTACCACAACCCCTACAAACGGCCAGCAGGGAATCTTTGGAAGAAGTGGTCCATGC

1022/6881
FIGURE 954B

TGATCAGAGCTCCTGCACTTTTGATAACCTGAGTCCCGGCTGGAGTACAATGTCAGTGTTCACACTGTCAAGGA
TGACAAGGAAAGTGTCCCTATCTCTGATACCATCATCCAGCTGTTCCCTCCCTCCCACTGACCTGCGATTACACAA
CATTGGTCCAGACACCATGCGTGTACCTGGGCTCCACCCCATCCATTGATTAAACCAACTTCCTGGTGCGTTA
CTCACCTGTGAAAAATGAGGAAGATGTTGCAGAGTTGTCAATTTCTCCTTCAGACAATGCAGTGGTCTTAACAAA
TCTCCTGCCTGGTACAGAATATGTAGTGAGTGTCTCCAGTGTCTACGAACAACATGAGAGCACACCTCTTAGAGG
AAGACAGAAAACAGGTCTTGATTCCCCAACTGGCATTGACTTTTCTGATATTACTGCCAACTCTTTTACTGTGCA
CTGGATTGCTCCTCGAGCCACCATCACTGGCTACAGGATCCGCCATCATCCCGAGCACTTCAGTGGGAGACCTCG
AGAAGATCGGGTGCCCCACTCTCGGAATTCCATCACCCCTACCAACCTCACTCCAGGCACAGAGTATGTGGTTCAG
CATCGTTGCTCTTAATGGCAGAGAGGAAAGTCCCTTATTGATTGGCCAACAATCAACAGTTTCTGATGTTCCGAG
GGACCTGGAAGTTGTTGCTGCGACCCCAACAGCCTACTGATCAGCTGGGATGCTCCTGCTGTACAGTGAGATA
TTACAGGATCACTTACGGAGAGACAGGAGGAAATAGCCCTGTCCAGGAGTTCACTGTGCCTGGGAGCAAGTCTAC
AGCTACCATCAGCGGCCCTTAAACCTGGAGTTGATTATACCATCACTGTGTATGCTGTCACTGGCCGTGGAGACAG
CCCCGCAAGCAGCAAGCCAATTTCCATTAAATTACCGAACAGAAATTGACAAACCATCCAGATGCAAGTGACCGA
TGTTTCAGGACAACAGCATTAGTGTCAAGTGGCTGCCCTCAAGTTCCCTGTTACTGGTTACAGAGTAACCAACAC
TCCCCAAAATGGACCAGGACCAACAAAACTAAACTGCAGGTCCAGATCAAACAGAAATGACTATTGAAGGCTT
GCAGCCACAGTGGAGTATGTGGTTAGTGTCTATGCTCAGAATCCAAGCGGAGAGAGTCAGCCTCTGGTTCAGAC
TGCAGTAACCAACATTGATCGCCCTAAAGGACTGGCATTCACTGATGTGGATGTCGATTCCATCAAAATTGCTTG
GGAAAGCCACAGGGGCAAGTTTCCAGGTACAGGGTGACCTACTCGAGCCCTGAGGATGGAATCCATGAGCTATT
CCCTGCACCTGATGGTGAAGAAGACACTGCAGAGCTGCAAGGCCTCAGACCGGGTCTGAGTACACAGTCAGTGT
GGTTGCCTTGCCAGATGATATGGAGAGCCAGCCCTGATTGGAACCCAGTCCACAGCTATTCTGCACCAACTGA
CCTGAAGTTCACTCAGGTACACCCACAAGCCTGAGCGCCAGTGGACACCAACCAATGTTTCAGCTCACTGGATA
TCGAGTGCGGGTGACCCCAAGGAGAAGACCGGACCAATGAAAGAAATCAACCTTGCTCCTGACAGCTCATCCGT
GGTTGTATCAGGACTTATGGTGGCCACCAATATGAAGTGAGTGTCTATGCTCTTAAGGACACTTTGACAAGCAG
ACCAGCTCAGGGAGTTGTACCACTCTGGAGAATGTCAGCCACCAAGAAGGGCTCGTGTGACAGATGCTACTGA
GACCACCATCACCATTAGCTGGAGAACCAAGACTGAGACGATCACTGGCTTCCAAGTTGATGCCGTTCCAGCCAA
TGGCCAGACTCCAATCCAGAGAACCATCAAGCCAGATGTCAGAAGCTACACCATCACAGGTTTACAACCAGGCAC
TGACTACAAGATCTACCTGTACACCTGAATGACAATGCTCGGAGCTCCCTGTGGTTCATCGACGCCCTCCACTGC
CATTGATGCACCATCCAACCTGCGTTTCTTGCCACCAACCAATTCCTTGCTGGTATCATGGCAGCCGCCACG
TGCCAGGATTACCGGCTACATCATCAAGTATGAGAAGCCTGGGTCTCCTCCCAGAGAAGTGGTCCCTCGGCCCGG
CCCTGGTGTACAGAGGCTACTATTACTGGCCTGGAACCGGGAACCGAATATACAATTTATGTCATTGCCCTGAA
GAATAATCAGAAGAGCGAGCCCTGATTGGAAGGAAAAAGACAGACGAGCTTCCCCAACTGGTAACCTTCCACA
CCCCAATCTTCATGGACCAGAGATCTTGGATGTTCTTCCACAGTTCAAAAGACCCCTTTCGTACCCACCCCTGG
GTATGACACTGGAATGGTATTACGCTTCTTGCACTTCTGGTCAGCAACCCAGTGTGGGCAACAAATGATCTT
TGAGGAACATGGTTTTAGGCGGACCACACCGCCACAACGGCCACCCCATAGGCATAGGCCAAGACCATAACC
GCCGAATGTAGGACAAGAAGCTCTCTCTCAGACAACCATCTCATGGGCCCCATTCCAGGACACTTCTGAGTACAT
CATTTTCATGTATCTCTGTTGGCACTGATGAAGAACCCTTACAGTTTCAGGGTCTCTGGAACCTTACCACTGCCAC
TCTGACAGGCCTCACCAGAGGTGCCACCTACAACGTCATAGTGGAGGCACTGAAAGACCAGCAGAGGCATAAGGT
TCGGGAAGAGGTTGTTACCGTGGGCAACTCTGTCAACGAAGGCTTGAACCAACCTACGGATGACTCGTGCTTTGA
CCCCACACAGTTTCCCATATGCCGTTGGAGATGAGTGGGAACGAATGTCTGAATCAGGCTTTAAACTGTTGTG
CCAGTGCTTAGGCTTTGGAAGTGGTCATTTCAGATGTGATTTCATCTAGATGGTGCCATGACAATGGTGTGAAC
CAAGATTGGAGAGAAGTGGGACCGTCAGGGAGAAAATGGCCAGATGATGAGCTGCACATGTCTTGGGAACGGAAA
AGGAGAATTCAAGTGTGACCCCTCATGAGGCAACGTGTTATGATGATGGGAAGACATACCACGTAGGAGAACAGTG
GCAGAAGGAATATCTCGGTGCCATTTGCTCCTGCACATGCTTTGGAGGCCAGCGGGCTGGCGCTGTGACAAC
CCGCAGACCTGGGGGTGAACCCAGTCCCGAAGGCACTACTGGCCAGTCTACAACCAAGTATTCTCAGAGATACCA
TCAGAGAACAAACACTAATGTTAATTGCCCAATTGAGTGCTTCATGCCTTTAGATGTACAGGTGACAGAGAAGA
TTCCCGAGAGTAAATCATCTTTCCAATCCAGAGGAACAAGCATGTCTCTCTGCCAAGATCCATCTAAACTGGAGT
GATGTTAGCAGACCCAGCTTAGAGTTCTTTCTTTCTTAAGCCCTTTGCTCTGGAGGAAGTCTCCAGCTTCA
GCTCAACTCACAGCTTCTCAAGCATCACCTGGGAGTTTCTGAGGGTTTTCTCATAAATGAGGGCTGCACATT

1023/6881
FIGURE 954C

GCCTGTTCTGCTTCGAAGTATTCAATACCGCTCAGTATTTTAAATGAAGTGATTCTAAGATTTGGTTTGGGATCA
ATAGGAAAGCATATGCAGCCAACCAAGATGCAAATGTTTTGAAATGATATGACCAAAATTTTAAGTAGGAAAGTC
ACCCAAACACTTCTGCTTTTCACTTAAGTGTCTGGCCCGCAATACTGTAGGAACAAGCATGATCTTGTACTGTGA
TATTTTAAATATCCACAGTACTCACTTTTTCCAAATGATCCTAGTAATTGCCTAGAAATATCTTCTCTTACCTG
TTATTTATCAATTTTCCAGTATTTTATACGGAAAAAATTGTATTGAAAACACTTAGTATGCAGTTGATAAGA
GGAATTTGGTATAATTATGGTGGGTGATTATTTTTTATCTGTATGTGCCAAAGCTTTACTACTGTGGAAAGACA
ACTGTTTTAATAAAAGATTTACATTCC

1024/6881
FIGURE 955

CGGCAGCCCTCCTACCTGCGCACGTGGTGCCGCTGCTGCTGCCTCCCGCTCGCCCTGAACCCAGTGCCTGCAGCC
ATGGCTCCCGGCCAGCTCGCCTTATTAGTGTCTCTGACAAAACCGGCCTTGTGGAATTGCAAGAAACCTGACC
GCTCTTGGTTTGAATCTGGTTCGCTTCCGGAGGGACTGCAAAAGCTCTCAGGGATGCTGGTCTGGCAGTCAGAGAT
GTCTCTGAGTTGACGGGATTTCTGAAATGTTGGGGGGACGTGTGAAAACCTTGCATCCTGCAGTCCATGCTGGA
ATCCTAGCTCGTAATATCCAGAAGATAATGCTGACATGGCCAGACTTGATTTCATCTTATAAGAGTTGTTGCC
TGCAATCTCTATCCCTTTGTAAAGACAGTGGCTTCTCCAGGTGTAACCTGTTGAGGAGGCTGTGGAGCAAATTGAC
ATTGGTGGAGTAACCTTACTGAGAGCTGCAGCCAAAACACGCTCGAGTGACAGTGGTGTGTGAACCAGAGGAC
TATGTGGTGGTGTCCACGGAGATGCAGAGCTCCGAGAGTAAGGACACCTCCTTGGAGACTAGACGCCAGTTAGCC
TTGAAGGCATTCACTCATACGGCACAATATGATGAAGCAATTTTCAAGATTATTTAGGAAACAGTACAGCAAAGGC
GTATCTCAGATGCCCTTGAGATATGGAATGAACCCACATCAGACCCCTGCCAGCTGTACACACTGCAGCCCAAG
CTTCCCATCACAGTTCTAAATGGAGCCCTGGATTTATAAACTTGTGCGATGCTTTGAACGCCCTGGCAGCTGGTG
AAGGAACTCAAGGAGGCTTTAGGTATTCCAGCCGCTGCCTCTTTCAAACATGTCAGCCCAGCAGGTGCTGCTGTT
GGAATTCCTCAGTGAAGATGAGGCCAAAGTCTGCATGGTTTATGATCTCTATAAAACCTCACACCCATCTCA
GCGGCATATGCAAGAGCAAGAGGGGCTGATAGGATGTCTTCAATTTGGTGATTTTGTGTCATTGTCCGATGTTTGT
GATGTACCAACTGCAAAAATTATTTCCAGAGAAGTATCTGATGGTATAATTGCCCCAGGATATGAAGAAGAAGCC
TTGACAATACTTTCCAAAAGAAAAATGGAACCTATTGTGTCCTTCAGATGGACCAATCTTACAAACCAGATGAA
AATGAAGTTCGAACCTCTCTTTGGTCTTCATTAAAGCCAGAAGAGAAAATAATGGTGTGCTCGACAAGTCATTATTT
AGCAATGTTGTTACCAAAAATAAAGATTTGCCAGAGTCTGCCCTCCGAGACCTCATCGTAGCCACCATGCTGTC
AAGTACACTCAGTCTAACTCTGTGTGCTACGCCAAGAACGGGCAGGTATCGGCATTGGAGCAGGACAGCAGTCT
CGTATACACTGCACTCGCCTTGCCAGGAGATAAGGCAAACTATTGGTGGCTTAGACACCATCCACAAGTGCTTTTCG
ATGAAGTTTAAACAGGAGTGAAGAGAGCAGAAATCTCCAATGCCATCGATCAATATGTGACTGGAACCATTTGGC
GAGGATGAAGATTTGATAAAGTGAAGGCACCTGTTGAGGAAGTCCCTGAGTTACTCACTGAGGCAGAGAAGAAG
GAATGGGTTGAGAACTGACTGAAGTTTCTATCAGCTCTGATGCCCTTCTTCCCTTTCCGAGATAACGTAGACAGA
GCTAAAAGGAGTGGTGTGGCGTACATTGCGGCTCCCTCCGGTTCTGCTGCTGACAAAGTTGTGATTGAGGCTGTC
GACGAACCTGGGAATCATCTCGCTCATACGAACCTTCGGCTCTTCCACCACTGATTTTTACCACACACTGTTTTTT
GGCTTGCTTATGTGTAGGTGAACAGTCACGCCTGAACTTTGAGGATAACTTTTTAAAAAATAAAACAGTATCT
CTTAAAACAATGTTTTGATCTACATAAACATTGTAAAAATTTTCAATCACGCTTTTTTAACCTTCTTACCACAAAA
AAATGATAAGTGGGTGAAGTGATGGTTATGTTAATTAGCGTGC

1025/6881
FIGURE 956

MAPGQLALFSVSDKTGLVEFARNLTALGLNLVASGGTAKALRDAGLAVRDVSELTGFPEMLGGRVKTLHPAVHAG
ILARNIPEDNADMARLDFNLIRVVACNLYPFVKTVASPGVTVEEAVEQIDIGGVTLRLAAAKNHARVTVVCEPED
YVVVSTEMQSSESKDTSLETRRQLALKAFTHTAQYDEAISDYFRKQYSKGVSQMPLRYGMNPHQTPAQLYTLQPK
LPITVLNGAPGFINLCDALNAWQLVKELKEALGIPAAASFKEHVSPAGAAVGIPLSEDEAKVCMVVDLYKTLTPIS
AAYARARGADRMSSFDFVALSDVCDVPTAKIISREVSDGIIAPGYEEEALTISSKKKNGNYCVLQMDQSYKPDE
NEVRTLFGLHLSQKRNNGVVDKSLFSNVVTKNKDLPESALRDLIVATIAVKYTSNSVCYAKNGQVIGIGAGQQS
RIHCTRLAGDKANYWVLRHHPQVLSMKFKTGKRAEISNAIDQYVTGTIGEDDLIKWKALFEEVPELLTEAEKK
EWVEKLTEVSISSDAFFPFRDNVDRAKRSKVAYIAAPSGSAADKVVEACDELGIILAHTNLRLFHH

1026/6881
FIGURE 957

GCAGCCAATTAAGCCAACTGAGTTCCTTTTCCTTGTGGGGGCCAGTGTGCAATGGCTGCACACAGCAGCTTCCTT
GGTAGTGTACACAGCCTGTTGGTTGTATGGGTTGCTCTGAGGGACCTTGGAGACAGGCCTTTCCAATGGATGTTT
ATGTTTCTGACCTTGCGCTACCCCAATGTAGGCTCCAAACAGGCATGCGAGGTGCCTTTGGAATGCCCCAGGGCA
CTGTGGCCAGGGTTACATTGGCCAAGTTATCATGTCCATCCGCACCAAGCTGCAGAACAAGGAGCATGTGATTG
AGGCCCTGCGCAGGGCCAAGTTCAAGCTTCCTGGCCACCAGAAGATCCACATCTCAAAGAAGTGGGGCTTCACCA
AGTTCAATGCTGATGAATTTGAAGACATGGTGGCTGAGAAGTGGCTCATCCCAGATGGCTGTGGGGTCAAGTACA
TCCCCAATCGTGGCCCTCTGGACAAGTGGCGGGCCCTGCACTCATGAGGGCTTCCAATGTGCTGCCCCCTCTTA
ATACTCACCAATAAATTCTACTTCCTGTCAAAAAAATGTCACTAATGTACCCTAAATTTTCTTTTTTATTTTAT
TTTTTATTTATTTATTTATTTTGTGAGATGGTGTCTTGCTCTTTTGCCAGACTGGAGTGAAGTAGCATGATCT
TGGCTTACTGCAACCTCCACCCCCAG

1027/6881
FIGURE 958

GGCGGGCGACCAAAGCGCCTGAGGACCGGCAACATGGTGCGGTCGGGGAATAAGGCAGCTGTTGTGCTGTGTATG
GACGTGGGCTTTACCATGAGTAACTCCATTCCCTGGTATAGAATCCCCATTTGAACAAGCAAAGAAGGTGATAACC
ATGTTTGTACAGCGCAGGTGTTTGTGAGAACAAAGGATGAGATTGCTTTAGTCCTGTTTGGTACAGATGGCACT
GACAATCCCCCTTCTGGTGGGGATCAGTATCAGAACATCACAGTGCACAGACATCTGATGCTACCAGATTTTGAT
TTGCTGGAGGACATTGAAAGCAAATCCAACCAGGTTCTCAACAGGCTGACTTCCTGGATGCACTAATCGTGAGC
ATGGATGTGATTCAACATGAAACAATAGGAAAGAAGTTTGAGAAGAGGCATATTGAAATATTTACTGACCTCAGC
AGCCGATTTCAGCAAAAGTCAGCTGGATATTATAATTCATAGCTTGAAGAAATGTGACATCTCCCTGCAATTTCTC
TTGCCTTTCTCACTTGGCAAGGAAGATGGAAGTGGGGACAGAGGAGATGGCCCCCTTCGCTTAGGTGGCCATGGG
CCTTCCTTTCCACTAAAAGGAATTACCGAACAGCAAAAAGAAGGTCTTGAGATAGTGAAAAATGGTGATGATATCT
TTAGAAGGTGAAGATGGGTTGGATGAAATTTATTTCATTTCAGTGAGAGTCTGAGAAAATGTGCGTCTTCAAGAAA
ATTGAGAGGCATTCCATTCACTGGCCCTGCCGACTGACCATTGGCTCCAATTTGTCTATAAGGATTGCAGCCTAT
AAATCGATTCTACAGGAGAGAGTTAAAAAGACTTGGACAGTTGTGGATGCAAAAACCTAAAAAAGAAGATATA
CAAAAAGAAACAGTTTATTGCTTAAATGATGATGATGAAACTGAAGTTTTAAAGAGGATATTATTCAAGGGTTC
CGCTATGGAAGTGATATAGTTCTTTCTCTAAAGTGGATGAGGAACAAATGAAATATAAATCGGAGGGGAAGTGC
TTCTCTGTTTTGGGATTTTGTAAATCTTCTCAGGTTTCAGAGAAGATTCTTCATGGGAAATCAAGTTCTAAAGGTC
TTTGACGAAGAGATGATGAGGCAGCTGCAGTTGCACTTTCTCCCTGATTTCATGCTTTGGATGACTTAGACAIG
GTGGCCATAGTTTCGATATGCTTATGACAAAAGAGCTAATCTCAAGTCGGCGTGGCTTTTCTCATATCAAGCAT
AACTATGAGTGTTTGTGTATGTGCAGCTGCCTTTTCATGGAAGACTTGCAGCAATACATGTTTTTCATCCTTGAAA
AACAGTAAGAAATATGCTCCCACCGAGGCACAGTTGAATGCTGTTGATGCTTTGATTGACTCCATGAGCTTGGA
AAGAAAGATGAGAAGACAGACACCCTTGAAGACTTGTTCACCAACCACAAAATCCCAAATCCTCGATTTTCAGAGA
TTATTTTCAGTGCTGCTGCACAGAGCTTTACATCCCCGGGAGCCTCTACCCCAATTTCAGCAGCATATTTGGAAT
ATGCTGAATCCTCCCGCTGAGGTGACAAACAAAAGTCAGATTCTCTCTCTAAAATAAAGACCCTTTTCTCTCTG
ATTGAAGCCAAGAAAAGGATCAAGTGACTGCTCAGGAAATTTTCCAAGACAACCATGAAGATGGACCTACAGCT
AAAAAATTAAGACTGAGCAAGGGGGAGCCCACTTCAGCGTCTCCAGTCTGGCTGAAGGCAGTGTCACCTCTGTT
GGAAGTGTAATCCTGCTGAAAACCTCCGTGTTCTAGTGAAACAGAAGAAGGCCAGCTTTGAGGAAGCGAGTAAC
CAGCTCATAAATCACATCGAACAGTTTTTGGATACTAATGAAACACCGTATTTTATGAAGAGCATAGACTGCATC
CGAGCCTTCCGGGAAGAAGCCATTAAAGTTTCAGAAGAGCAGCGCTTTAACAACTTCTGAAAGCCCTTCAAGAG
AAAGTGAAATTAACAATTAATCATTCTGGGAAATTTGTGTCCAGGATGGAATTACTCTGATCACCAAGAG
GAAGCCTCTGGAAGTTCTGTACAGCTGAGGAAGCCAAAAGTTTCTGGCCCCAAGACAAAACCAAGTGAGAC
ACAGCAGCTGTATTTGAAGAAGGTGGTGATGTGGACGATTTATTGGACATGATATAGGTCGTGGATGATGGGGA
ATCTAAGAGAGCTGCCATCGCTGTGATGCTGGGAGTTCTAACAAAACAAGTTGGATGCGGCCATTCAAGGGGAGC
CAAAATCTCAAGAAATTCAGCAGGTTACCTGGAGGCGGATCATCTAATTCCTGTGGAATGAATACACACATA
TATATTACAAGGGATAATTTAGACCCCATACAAGTTTATAAAGAGTCATTGTTATTTTCTGGTTGGTGATTATT
TTTTCTGTGGTCTTACTGATCTTTGTATATTACATACATGCTTTGAAGTTTCTGGAAAGTAGATCTTTTCTTGAC
CTAGTATATCAGTGACAGTTGCAGCCCTTGTGATGTGATTAGTGCTCATGTGGAACCATGGCATGGTTATTGAT
GAGTTTCTTAACCCTTTCCAGAGTCCTCCTTTGCCTGATCCCTCAACAGCTGTCAACACTTGTGTTGAGCAAGCA
GTAGCATTGCTTCTCCCAACAAGCAGCTGGGTTAGGAAAACCATGGGTAAGGACGGACTCACTTCTCTTTTTTA
GTTGAGGCCTTCTAGTTACCACATTACTCTGCCTCTGTATATAGGTGGTTTTCTTTAAGTGGGGTGGGAAGGGGA
GCACAATTTCCCTTCATACTCCTTTTAAGCAGTGAGTTATGGTGGTGGTCTCATGAAGAAAAGACCTTTTGCCCC
AATCTCTGCCATATCAGTGAACCTTTAGAACTCAAAAAGTGAAGAAATTTACTTCAGTAGTTAGAATTATATCAC
TTCATCTGTCTCTACTTGCAAGCCTCAAAGAGAGAAAGTTTCGTTATATTAAACACTTAGGTAACCTTTTCGGTC
TTTCCCATTTCTACCTAAGTCAGCTTTTCATCTTTGTGGATGGTGTCTCCTTTACTAAAATAAGAAAATAACAAAGC
CCTTATTCTCTTTTTTCTGTCTCATTCTTGCTTGAAGTTCCAGTTTCTCTTTGGTGACAGACTTCTTGTA
CCAGTCACCTCTGTCTTCAGCACCCCTCATAAGTCGTCACTAATACACAGTTTTGTACATGTAACATTAAAGGCA
TAAATGACTC

1028/6881
FIGURE 959

MVRSGNKAADVLCMDVGFTMSNSIPGIESPFQAKKVITMFVQRQVFAENKDEIALVLFGTDGTDNPLSGGDQYQ
NITVHRHMLPDPFDLLEDIESKIQPGSQADFLDALIVSMDVIQHETIGKKFEKRHIEIFTDLSSRFSKSQLDII
IHSLKKCDISLQFFLPFSLGKEDGSGDRGDGPFRLGGHGPSFPLKGITEQQKEGLEIVKMVMISLEGEDGLDEIY
SFSESLRKLCVFKKIERHSIHWPCRLTIGSNLSIRIAAYKSILQERVKKTWTVVDAKTLKKEDIQKETVYCLNDD
DETEVLKEDIIQGFYRGSDIVPFSKVDEEQMKYKSEGKCFSVLGFCKSSQVQRFFMGNQVLKVFAARDDEAAAV
ALSSLIHALDDLDMAIVRYAYDKRANPQGVAFPHIKHNYECLVYVQLPFMEDLRQYMFSSLKNSKKYAPTEAQ
LNAVDALIDSMSLAKKDEKDTLEDLPPTTKIPNPRFQRLFQCLLHRALHPREPLPPIQQHIWNMLNPPAEVTTK
SQIPLSKIKTLFPLIEAKKKDQVTAQEIFQDNHEDGPTAKKLKTEQGGAHFSVSSLAEGSVTSVGSVNPAENFRV
LVKQKKASFEEASNQLINHIEQFLDTNETPYFMKSIDCIRAFREEAIKFSEEQRFNFLKALQEKVEIKQLNHFV
EIVVQDGITLITKEEASGSSVTAEAEAKKFLAPKDKPSGDTAAVFEEGGDVDDLLDMI

1029/6881
FIGURE 960A

CTTTTTGAGGCAAGTTGGATGAACAGAGAAGGGAAGAGAGGAAGAACGAGAGGAAGAGAAGGGAAGGAAGTGTTT
GTGTAGAAGAGAGAGAAAGACGAATAGAGTTAGGAAAAGGAAGACAAGCAGGTGGGCAGGAAGGACATGCACCGA
GACCAGGCAGGGGCCCAACTTTACGTCCAGCCCTGGCCTGGGGTCGGGAGAGGTGGGCGCTAGAAGATGCAGCC
CAGGATGTGGCAATCAATGACACTATTGGGGTTTCCCAGGATGGATTGGTCAGGGGGAGAAAGGAAAGGCAAAA
CACTCCAGGACCTCTCCCGATCTGTCTCCTCCTCTAGCCAGCAGTATGGACAGCTGGACCCCTGAACTTCCTCT
CCTCTTACCTGGGCAGAGTGTGTCTCTCCCAAAATTTATAAAAACATAAATGCATTCCATTCTCTGAAAGCAA
AACAAATTCATAATTGAGTGATATTAATAGAGAGGTTTTTCGGAAGCAGATCTGTGAATATGAAATACATGTGCA
TATTTCAATTCCTCCAGGCAGACATTTTTTAGAAATCAATACATGCCCAATATTGGAAAGACTTGTTCTTCCACGG
TGACTACAGTACATGCTGAAGCGTGCCGTTTCAGCCCTCATTAAATCAATTTGTAAGTAGCGCAGCAGCCCTCTG
TGGGGGAGGATAGGCTGAAAAAAAAGTGGGCTCGTATTTATCTACAGGACTCCATATAGTCATATATAGGCAT
ATAAATCTATTCTTTTTCTTTGTTTTTTCTTTCTTCTTTCTTTCAAAGGTTTGCAATTAACCTTTCAAAGTAGT
TCCTATAGGGGCAATTGAGGAGCTTCCTCATTCTGGGAAAACCTGAGAAAACCCATATTCTCCTAATACAACCCGTA
ATAGCATTTTTGCCTGCCTCGAGGCAGAGTTTTCCCGTGAGCAATAAACTCAGCTTTTTTGTTGGGGCACAGTACTG
GATTTGACAGTGATTCCCCACGTGTGTTTCATCTGACCCACCGAGCCAGGCAGAGGCCAGCCCTCCGTGGTGAC
ACAGCACGCGCTCAGTCCATCCCATTTTAGTCTTTAAACCCCTCAGGAAGTCACAGTCTCCGGACACCACACCAC
ATGAGCCCCAACAGGTCCACGATGGATCCACCAGTCCCACCCAGCCTTTTCTTTTCATCTGAACAGAATGTGCAT
TTTTGGAAGCTTCCCTCACTCTCCATGCTGGCAGAGCAGGAGGGAGACTGAAGTAAGAGATGGCAGAGGGAGATG
GTGGCAAAAAGGTTTAGATGCAGGAGAACAGTAAGATGGATGGTTCCGGCCAGAGTCGATGTGGGGAGGAACAGA
GGGCTGAAGGGAGAGGGGGCTGACTGTTCCATTCTAGCTTTGGCACAAAGCAGCAGAAAGGGGGAAAAGCCAATA
GAAATTTCTTAGCTTCCCCACCATATGTATTTTCTAGGATTTGAGAGGAAAGAGAGGAAAATGGGGGAATGGGT
TGCAAAATAGAAATGAGCTTAATCCAGGCCGAGAGCCAGGGAAGGTGAGTAACCTTTAGGAGGGTGCTAGACTTT
AGAAGCCAGATAGGAAGAATCAGTCTAAACTGGCCATGCTTTGGAAGGGACAAGACTATGTGCTCCGCTGCCAC
CTTCAGCCTGCAATGAGGGACTGAGGCCACGAGTCTTCCAGCTCTTCTCCTCATTCTGGCCAGTCCCTGCATCC
TCCCTGGGGTGGAGGATGGAAGGAAAGCTGGGACAAGCAGGGAACGCATGATTACAGGATGCTGTCTACTCGGCAG
CCAGATTCCGAAACTCCCATTTCTCAATGACTTCTCAACCAATGGGTGGCCTTGCTGACTGTTCTTTAAGGCTGA
AGATATCCAGGAAAGGGGGCTTGACACTGGCCAAGGAGACCCCTTCGTGCTGTGGACACAGCTCTCTTCACTCT
TTGCTCATGGCATGACACAGCGGAGACCGCCTCCAACAACGAATTTGGGGCTACGAAGAGGAATAGCGAAAAAGC
AAATCTGTTTCAACTGATGGGAACCCATAGCTATAGAACTTGGGGGCTATCTCCTATGCCCTGGACAGGACAG
TTGGCTGGGGACAGGAGAAGTGCTCAATCTTCATGAGACAAAGGGGCCGATAGGGCCAGCAGCCACAAGGCCTT
GACCTGCCGAGTCAGCATGCCCATCTCTCTGCACAGCTGTCCCTAAACCCAACTCACGTTTCTGTATGTCTTA
GGCCAGTATCCCAAACCTCTTCCACGTCACTGTTCTTTCCACCCATTCTCCCTTTGCATCTTGAGCAGTTATCCA
ACTAGGATCTGCCAAGTGGATACTGGGGTGCCACTCCCTGAGAAAAGACTGAGCCAGGAATACAAGCTCCCC
CACATTCTCCAGCCTGGACCTAATCTTGAGAGGGCTCTCTCTTCACGACTGTGTCTGACTTTGAGCAGG
CTTCTGCCCTTGCGTTGGCTCTTTGCTGCCAGCCATCAGGTGGGGGATTAGAGCCTGGTGTAAAGTGCGCCAGAC
TCTTCCGGTTTCCAAAGTTTCGTGCCGGAACCCAAACCTGTGAGTCTCTTCTGCATGCAGGAGTTTCTCCTGGG
CAGCTGGTCACTCCCCAGAGAAGCTGGGCCTTCATGGACACATGGAACATAAGCCTCCCAATGGGAGTTCTGGCT
GAGCCCAGGTGGGGAGATCCTGGGAAGGGAGGCACTGGAGGAAGACGGCACCTCTTCCCCATGGCAGGGGTG
AGGGAGGCAGGTTTGAATGGTGCGAGTATGGCAATCTAAGCAGGGGTCTGGTCTCTTTGACTCCAGGCTGGCCT
TTGGCCGACTGTCTGCTCACCCAGAGACCTTGACTCCGACTATCCATGGCTCCGAATCTAAGTGCTGCCACT
CCCATGCTCACACCCACAGAAGGTCTTCCATCCCTTTAGATTGCTGCCCTCACTCCACAGTGAGGAAGATGCC
TCTGTCTTTCCACGACTGCCAGGAGATAGGGAAGCCAGCCAGGACTGACCTCCTTCTCCAGCCTGCCCTGA
CCCACCTGGCAAAGCAGGGCACATGGGGAGGAAGAGACTGGAACCTTTCTTGACAGCCAGGCCCTAGACAGACAG
GCCTGGGGACACTGGCCCCATGAGGGGAGGAAGGCAGGCGCACGAGGTCCAGGGAGGCCCTTTCTGATCATGCC
CCTTCTCTCCACCCCATCTCCCCACCACCTCTGTGGCCTCCATGGTACCCCCACAGGGCTGGCCTCCCCCTA
GAGGGTGGGCTCAACCACTGCTCCCGCCACGACCGGTAGTGAGACAGGGCTCCGCCCCAGAGATGGGAACCTCA
CCCCTCAAGGTGGGACAGTACCCCGGACCCATCCACTCACTCCTGAGAGGGCTCCGCCCCAGAGATGGGAACCTCA
GAGAAGAGCTCTAAGGAGAAGAAACCCCATAGCGTCAGAGAGGATATGTCTGGCTTCCAAGAGAAAGGAGGCTCC
GTTTTGCAAAGTGAGGAGGGACGAGGGACAGGGGTTTACCAGCCAGCAACCTGGGCCTTGTAAGTGTCTGTGT

1030/6881

FIGURE 960B

TTTAAAACCACTAAAGTGCAAGAATTACATTGCACTGTTTCTCCACTTTTTATTTTCTCTTAGGCITTTTGTCTTCT
ATTTCAAACATACTTTCTTGGTTTTCTAATGGAGTATATAGTTTAGTCATTTACAGACTCTGGCCTCCTCTCCT
GAAATCCTTTTGGATGGGGAAAGGGAAGGTGGGGAGGGTCCGAGGGGAAGGGGACCCAGCTTCCCTGTGCCCCG
TCACCCCACTCCACCAGTCCCCGGTCGCCAGCCGGAGTCTCCTCTCTACCGCCACTGTACACCCGTAGCCACAT
GGATAGCACAGTTGTCAGACAAGATTCCCTCAGATTCCGAGTTGCCTACCGGTGTTTTTCGTTGTTGTTGTTGTT
GTTTTCTTTTTCTTTTTTTTTTTGAAGACAGCAATAACCACAGTACATATTACTGTAGTTCTCTATAGTTTTAC
ATACATTCATACCATAACTCTGTTCTCTCCTCTTTTTTGTGTTTCACTTTAAAAACAAAAATAAACGATGATAAT
CTTTACTGGTGAAAAGGATGGAAAAATAAATCAACAAATGCAACCAGTTTGTGAG

1031/6881
FIGURE 961

MLWKGQDYVLRCPPSACNEGLRPTSLSSSSSILASPCILPGVEDGRKAGTSRERMIQGCCHSAARFRNSHSPMTS
STNGWPCDCSLRLKISRKGGLDTGQGDPFVLWTQLSSLFAHGMTQRRPPPTTNLGLRRGIAKKQICFN

1032/6881
FIGURE 962

CCTCTGTGGCCTCCATGGTACCCCCACAGGGCTGGCCTCCCCTAGAGGGTGGGCCTCAACCACCTGCTCCCGCCA
CGCACCGGTTAGTGAGACAGGGCTGCCACGGCAACCGCCAAGCCCCCTCAAGGTGGGACAGTACCCCGGACCCA
TCCACTCACTCCTGAGAGGGCTCCGGCCCAGAATGGGAACCTCAGAGAAGAGCTCTAAGGAGAAGAAACCCATA
GCGTCAGAGAGGATATGTCTGGCTTCCAAGAGAAAGGAGGCTCCGTTTTGCAAAGTGGAGGAGGGACGAGGGACA
GGGGTTTCACCAGCCAGCAACCTGGGCCTTGTACTGTCTGTGTTTTTAAAACCACTAAAGTGCAAGAATTACATT
GCACTGTTTCTCCACTTTTTATTTTCTCTTAGGCTTTTGTCTATTTCAAACATACTTTCTTGGTTTTCTAATG
GAGTATATAGTTTAGTCATTTACAGACTCTGGCCTCCTCTCCTGAAATCCTTTTGGATGGGGAAAGGGAAGGTG
GGGAGGGTCCGAGGGGAAGGGGACCCAGCTTCCCTGTGCCCGCTCACCCCACTCCACCAGTCCCGGTGCGCCAG
CCGGAGTCTCCTCTCTACCGCCACTGTTCACACCGTAGCCACATGGATAGCACAGTTGTCAGACAAGATTCCCTC
AGATTCCGAGTTGCCTACCGGTTGTTTTCGTTGTTGTTGTTGTTGTTTTTCTTTTCTTTTTTTTTTTGAAGACA
GCAATAACCACAGTACATATTACTGTAGTTCTCTATAGTTTACATACATTCATACCATAACTCTGTTCTCTCCT
CTTTTTGTTTTCACTTTAAAAACAAAAATAAACGATGATAATCTTTACTGGTGAAAAGGATGGAAAAATAAAT
CAACAAATGCAACCAGTTTGTGAGAAAAAAAAAAAAAGCCGAAAAAAAAAAAAA

1033/6881
FIGURE 963

MVFPQGWPPLEGGPQPPAPATHRLVRQGCHGNRQAPLKVGPYGPPIHSLRLRGLRPRMGTSEKSSKEKKPHSVRED
MSGFQEKGGSVLQSGGGTRDRGFTSQQPGPCTVCVFKTKVQELHCTVSPLFIFS

1034/6881
FIGURE 964

GGCGAGGGAGGAGGAAGAAGCGGAGGAGGCGGCTCCCGCGCTCGCAGGGCCGTGCCACCTGCCCCGCCCGCCGCT
CGCTCGCTCGCCCGCCGCGCCGCGCTGCCGACCGCCAGCATGCTGCCGAGAGTGGGCTGCCCCGCGCTGCCGCTG
CCGCCCGCCCGCTGCTGCCGCTGCTGCTGCTGCTACTGGGCGCGAGTGGCGGGCGGGCGGGGCGCGCGCGGAG
GTGCTGTTCCGCTGCCCCGCCCTGCACACCCGAGCGCTGGCCGCTGCGGGCCCCCGCCGGTTGCGCCGCCCGCC
GCGGTGGCCGAGTGGCCGGAGGCGCCCGCATGCCATGCGCGGAGCTCGTCCGGGAGCCGGGCTGCGGCTGCTGC
TCGGTGTGCGCCCGGCTGGAGGGCGAGGCGTGCGGCGTCTACACCCGCGCTGCGGCCAGGGGCTGCGCTGCTAT
CCCCACCCGGGCTCCGAGCTGCCCTGTCAGGCGCTGGTTCATGGGCGAGGGCACTTGTGAGAAGCGCCGGGACGCC
GAGTATGGCGCCAGCCCGGAGCAGGTTGCAGACAATGGCGATGACCACTCAGAAGGAGGCCTGGTGGAGAACCAC
GTGGACAGCACCATGAACATGTTGGCGGGGGAGGAGTGTGCGCCGGAAGCCCCTCAAGTCGGGTATGAAGGAG
CTGGCCGTGTTCCGGGAGAAGGTCACTGAGCAGCACCGGCAGATGGGCAAGGGTGGCAAGCATCACCTTGGCCTG
GAGGAGCCCAAGAAGCTGCGACCACCCCTGCCAGGACTCCCTGCCAACAGGAAGTGGACCAGGTCTTGGAGCGG
ATCTCCACCATGCGCCTTCCGGATGAGCGGGGCCCTCTGGAGCACCTCTACTCCCTGCACATCCCCAACTGTGAC
AAGCATGGCCTGTACAACCTCAAACAGTGCAAGATGTCTCTGAACGGGCAGCGTGGGGAGTGTGCTGGTGTGTGAAC
CCCAACACCGGAAGCTGATCCAGGGAGCCCCCACCATCCGGGGGACCCCGAGTGTATCTCTTCTACAATGAG
CAGCAGGAGGCTCGCGGGGTGCACACCCAGCGGATGCAGTAGACCGCAGCCAGCCGGTGCCTGGCGCCCTTGCCC
CCGCCCCCTCTCCAAACACCGGCAGAAAACGGAGAGTGCTTGGGTGGTGGGTGCTGGAGGATTTCCAGTTCTGA
CACACGTATTTATATTTGGAAAGAGACCAGCACCGAGCTCGGCACCTCCCCGGCCTCTCTCTTCCAGCTGCAGA
TGCCACACCTGCTCCTTCTTTGCTTTCCCCGGGGGAGGAAGGGGTTGTGGTGGGGAGCTGGGGTACAGGTTTGG
GGAGGGGGAAGAGAAATTTTATTTTGAACCCCTGTGTCCCTTTTGATAAGATTAAAGGAAGGAAAAGT

1035/6881
FIGURE 965

AGCGGATCTCCACCATGCGCCTTCCGGATGAGCGGGGCCCTCTGGAGCACCTCTACTCCCTGCACATCCCCAACT
GTGACAAGCATGGCCTGTACAACCTCAAACAGTGCTGGTGTGTGAACCCCAACACCGGGAAGCTGATCCAGGGAG
CCCCACCATCCGGGGGGACCCCGAGTGTATCTCTTCTACAATGAGCAGCAGGAGGCTCGCGGGGTGCACACCC
AGCGGATGCAGTAGACCGCAGCCAGCCGGTGCCTGGCGCCCTGCCCCCCGCCCCCTCTCCAAACACCGGCAGAAA
ACGGAGAGTGCTTGGGTGGTGGGTGCTGGAGGATTTTCCAGTTCTGACACACGTATTTATATTGGAAAGAGACC
AGCACCGAGCTCGGCACCTCCCCGGCCTCTCTCTTCCCAGCTGCAGATGCCACACCTGCTCCTTCTTGCTTTCCC
CGGGGAGGAAGGGGTTGTGGTTCGGGAGCTGGGGTACAGGTTTGGGGAGGGGAAGAGAAATTTTATTTTTG
AACCCTGTGTCCCTTTGCATAAGATTAAAGGAAGG

1036/6881
FIGURE 966

CTTCCTTTCTGGGCTCGGACCTAGGTCGCGGCGACATGGCCAAACGTACCAAGAAAGTCGGGATCGTCGGTAAAT
ACGGGACCCGCTATGGGGCCTCCCTCCGGAAAATGGTGAAGAAAATTGAAATCAGCCAGCACGCCAAGTACACTT
GCTCTTTCTGTGGCAAACCAAGATGAAGAGACGAGCTGTGGGGATCTGGCACTGTGGTTCCTGCATGAAGACAG
TGGCTGGCGGTGCCTGGACGTACAATACCACTTCCGCTGTACGGTAAAGTCCGCCATCAGAAGACTGAAGGAGT
TGAAAGACCAGTAGACGCTCCTCTACTCTTTGAGACATCACTGGCCTATAATAAATGGGTAAATTTATGTAACAA
AAAAAAAAAAAAAAAAA

1037/6881
FIGURE 967

MAKRIKKVGIVGKYGTRYGASLRKMKVKKIEISQHAQYTCSECGKTKMKRRAVGIWHCGSCMKTVAGGAWTYNTTS
AVTVKSAIRRLKELKDQ

1038/6881
FIGURE 968

CCCCCGCCACCTCACACACCCCATCCGGGAGACCCACGTGCAGCTGGGCCCCGGGCTGGCGCCGCACCTCGGGG
TCTGCGCGTCTCCCCGGCGCGTTCCGGGCGCAGCCCCACGACACGGACCCCTGGCGGGGCGGCATCCGAGAAGTC
GCAGGCGGAGAGGGGGCGTCTTGGAGGCCGGGCGGGGACGCTGCGGCCCGCGGCCACACAAAGGAGGCGGCGG
GAAGGCGGGGCAAGGCGGGCCGGGGGCGGGGCGGCAGGAAGGGGCGGGGGCCCCGCGCGGCGCCGCCGATAAAGC
CCCCGCGCGCGCGGCAGCCAGCTTGCCTGTGGGGCTGCCCGGGCTGCGCGGCGTCTGCAGGCGCCACCGCTGCC
TCTTTCCGGCTGTGACCTCCTCGCCGCCCGCGCTTGGCTGCGTCTCCGACTCCCCGCGCCGCCGAGACCAGGC
TCCCGCTCCGGTTGCGGCCGCACCGCCCTCCGCGGCCGCCCTTGGGGATCCAGCGAGCGCGGTCTCTTTGGTG
GAAGGAACCATGAACTGGCATCTCCCCCTCTTCTCTTGGCCTCTGTGACGCTGCCCTTCCATCTGCTCCCACTTC
AATCCTCTGTCTCTCGAGGAAGTGGCTCCAACACGGGGATCCAGGTTTTCAATCAGATTGTGAAGTCGAGGCCT
CATGACAACATCGTGATCTCTCCCCATGGGATTGCGTGGTCTTGGGGATGCTTCAGCTGGGGGCGGACGGCAGG
ACCAAGAAGCAGCTCGCCATGGTGATGAGATACGGCGTAAATGGAGTTGGTAAATATTAAAGAAGATCAACAAG
GCCATCGTCTCCAAGAAGAATAAAGACATTGTGACAGTGGCTAACGCCGTGTTTGTAAAGAATGCCTCTGAAATT
GAAGTGCCCTTTTGTACAGGAACAAAGATGTGTTCCAGTGTGAGGTCCGGAATGTGAACCTTTGAGGATCCAGCC
TCTGCCTGTGATTCCATCAATGCATGGGTTAAAAATGAAACCAGGGATATGATTGACAATCTGCTGTCCCCAGAT
CTTATTGATGGTGTGCTCACCAGACTGGTCTCGTCAACGCAGTGTATTTCAAGGGTCTGTGGAATCACCAGTTC
CAACCCGAGAACACAAAGAAACGCACCTTCGTGGCAGCCGACGGGAAATCCTATCAAGTGCCAATGCTGGCCCCAG
CTCTCCGTGTTCCGGTGTGGGTGCGACAAGTGCCCCCAATGATTATGGTACAACCTTCATTGAAGTGCCTACCAC
GGGGAAAGCATCAGCATGCTGATTGCACTGCCGACTGAGAGCTCCACTCCGCTGTCTGCCATCATCCACACATC
AGCACCAAGACCATAGACAGCTGGATGAGCATCATGGTGCCCAAGAGGGTGCAGGTGATCCTGCCCAAGTTCACA
GCTGTAGCACAAACAGATTGAAGGAGCCGCTGAAAGTTCTTGGCATTACTGACATGTTTGATTTCATCAAAGGCA
AATTTTGCAAAAATAACAAGGTGAGAAAACCTCCATGTTTCTCATATCTTGCAAAAAGCAAAAATTGAAGTCAGT
GAAGATGGAACCAAAGCTTCAGCAGCAACAACCTGCAATTCTCATTGCAAGATCATCGCCTCCCTGGTTTATAGTA
GACAGACCTTTTCTGTTTTTCATCCGACATAATCCTACAGGTGCTGTGTTATTTCATGGGGCAGATAAAACAAACC
TGAAGAGTATACAAAAGAAACCATGCAAAGCAACGACTACTTTGCTACGAAGAAAGACTCCTTTCTGCACTTTT
CATAGTTCTGTAAATATTTTTGTACATCGCTTCTTTTTCAAACTAGTTCTTAGGAACAGACTCGATGCAAGTG
TTTCTGTTCTGGGAGGTATTGGAGGGAAAAACAAGCAGGATGGCTGGAACACTGTACTGAGGAATGAATAGAAA
GGCTTCCAGATGTCTAAAGATTCTTTAACTACTGAACTGTTACCTAGGTAAACAACCTGTTGAGTATTTGCT
GTTTGTCAGTTCAGGAATTTTTGTTTTGTTTTGTCTATATGTGCGGCTTTTCAGAAGAAATTTAATCAGTGTGA
CAGAAAAAAAATGTTTTATGGTAGCTTTTACTTTTTATGAAAAAAAATTTATTTGCCTTTTAAATCTTTTCCC
CCATCCCCCTCCAAAGTCTTGATAGCAAGCGTTATTTTGGTGGTAGAAACGGTGAAATCTCTAGCCTCTTTGTGT
TTTTGTGTTGTTGTTGTTGTTGTTTATATAATGCATGTATTCACTAAAATAAAATTTAAAAAATCCTGTCTT
GCTAGACAAGGTTGCTGTTGTGCAGTGTGCCTGTCACTACTGGTCTGTACTCCTTGGATTGCAATTTTGTATTT
TGTACAAAGTAAAAATAAACTGTTATGAGTAGT

1039/6881
FIGURE 969

MNWHLPFLLLASVTLP SICS HFNPLSLEELGSNTGIQVFNQIVKSRPHDNIVISPHGIASVLGMLQLGADGRTKK
QLAMVMRYGVNGVGKILKKINKAIVSKKNKDIVTVANAVFVKNASEIEVPFVTRNKDVFQCEVRNVNFEDPASAC
DSINAWVKNETRDMIDNLLSPDLIDGVLTRLVLVNAVYFKGLWKS RFQ PENTKKRTFVAADGKSYQVPMLAQLSV
FRCGSTAPNDLWYNFIELPYHGESISMLIALPTESSTPLSAIIPHISTKTIDSWMSIMVEKRVQVILPKFTAVA
QTDLKEPLKVLGITDMFDSSKANFAKITRSENLVSHILQKAKIEVSEDGTKASAATTAILIARSSPPWFIVDRE
FLFFIRHNPTGAVLFMGQINKP

1040/6881
FIGURE 970

ATGCGTGCTGAAAAGAGAAAAAAGAATGCCCCAGAGGAGGTCAGCAGGCTTAAAAGTATTCTCAAAGTAGACGAT
GACGTTTTAATGAAAGATGTTCAAGAGATAGCAACTGTGGTGGTACCCAAACCCAAACATTGCCAAGAGAAAATG
CAATGTGAGGTTAAAGATGAAAAAGATGACATGAAAATGGAGACTGATATTAAGAGAAACAAAAAGACTCTTCTA
GACCAGCATGGACAGTACCCAATATGGATGAACCAAGGCAAAGAAAAAGGCTAAAGGCAAAGCGAGAGAAAAGA
AAGGAGAAAAACAAAGCAAAAGCAGTGAAAGTGGCAAAGGGTTTGGCCTGTAGTATTGTTTCCATAATAGTCCAT
TTCGCAGAAATAAGAAGGAGAATAAACTTAGAGAAAAAGAGGAGGCCAAAAGGGGCAAGGTAGTGGCTACTTGC
AGAAACCTGAAAAATATGTGTGACAAAGCCGCCATAATAAGTGGCAGCCGAGCTGCAGCAGCCCGTCTCAGGGGC
ACCGCAGCCTCCCCGGGCCCTACGGCCGCCGCCAGCAGGATGGCTGGAATGGCCTTAGTCATGAGGCTTTTAGA
ATTGTTTCAAGGCAGGATTATGCGTCAGAAGCAATCAATGGAGCAGCTGGTGGTGTATTATTTGGGTACTACCAAC
TCCTGTGTGGCAGTTATGGAAGGTAACAAGCAAAGGTGCTGGAGAATGCCGAAGGTGCCAGAAGTACCCCTTCA
GTTGTGGCCTTTACAGCAGATGTACAGAAAGACATTAAAAATATTCCTTTTAAATTTGTCTGTGCCTCCAATGGT
GATGCCTGGGTTGAGGCTCATGGGAACTGTATTCTCCAAGTCAGATCGGAGCATTGTGTGCGATGAAGATGAAA
GAGACTGCAGAAAATTACTTGGGGCACACAGCAAAAAATGCTGTGATCGCAGTCCCAGCTTATTTCAATGACTTG
CAGAGGCAGGCCACTAAAGATGCTGGCCAGATATCTGGACTGAATGTGCTTCGGGTGATTAATGAACCCACAGCT
GCTGCTCTTGCCATGTTCTAGACAAATCAAAGACAAAGTCATTGCTGTATATGATTTAGGTGGTGGAACTTTT
GATATTTCTATCCTGGAATTTAGAAAGGAGTATTCGAGGTGAAATCCACAAATGGGGACACTTTCTTAGGTGGG
GAAGACTTTGACCAGGCCTTGCTACAGCACATTGTGAAGGAGTTCAAGAGAGAGACGGGGGTGATTTGACCAAA
GACAACGTGGCATTTCAGAGGGTGTGGGAAGCTGCTGAAAAGGCTAAATGTGAACTCTCCTCATCTGTGCAGACT
GACATCAATTTGCCCTATCTTACAATGGATTCTTCTGGACCCAAGCATTGGAATACAAAGTTGACCCATGCTCAA
TTTGAAGGGATTGCTACTGATCTAATCAGGAGGACTATCGCTCCATGCCAAAAAGCTATGCAAGATGCAGAACTC
AGCAAGAGTGATATAGGAGAAGTGATTCTTGTGGAGCCCCCAAGTAAAGCTGTCAATCCTGATGAGGCTGTGGCC
ATTGGAGCTGCCATTGAGGGAGATGTGTTGGCCGGCGATGTACAGATGTGCTGCTCCTTGATGTCATTCCCTG
TCTCTGGGTATTGAACTCTGCGAGGTGTCTTTACCAAACCTTATTAATAGGAATACCACTATTTCAACCAAGAAG
AGCCAGGTATTCTCAACTGCTGCTGATGGGCAGACACAAGTGGAAATTAAAGTGTGTCAGGGTGAGAGAGAGATG
GCTGGAGACAACAACTCCTTGACAGTTTACTTTGATTGGAATTCCACCAGCCCCCTCGTGGAGTTCTCTCAGATC
AAAGTTACATTTGACATTGATGCCAATGGGATCATACATGTTTCAGCTAAAGATAAAGGCACAGGACATGAGCAG
CAGATTGTAATCCAGTCTTCTGGTGGGTTAAGCAAAGATGATGTTGAAAATATGGTTAAAAATGCAGAGAAATAT
GCTAAGGAAGACCAGCGAAGGAAGGAATGA

1041/6881
FIGURE 971

CGGAGCTGTCCATCAGCACCAAAGGCCGCGGGCGGGCTCAGGGCATGGGGCCGCGGTTCTGGGGCGGGCCGAGCC
CCGGCTCCTGCGCCTTCCCCCTTCTCAGGCCCAGCCCCAGTTCCCGGACGCCGCGGGACTGGAGTGCCAGCCGGT
GTTGGACGTGGAGCGGCGCCGCCACC GCGCCGACACCATTCTCTCCGGCCCAGCAGCCCCCTTCTCGCACGACG
GACTTTCCCTGGACCCCACTCAGTTGGAGCCTCTGGCGCCCCGAACCCCGGCCCTCGGGCCTCTGCACAGCCT
CTTTCACTCAGAAGCTCAGGTCGCCTCCAGCCCAGCACTATGCCGGGGACTGTGGCAACACTGCGGTTCCAGCTG
CTGCCCCCTGAGCCAGATGATGCCTTCTGGGGTGCACCTTGTGAACAGCCCCCTGGAGCGCAGGTACCAGGCACTG
CCGGCCCTCGTCTGCATCATGTGCTGTTTGTGGAGTCGTCTACTGCTTCTTCGGTTACCGCTGCTTCAAGGCA
GTGCTCTTTCTACTGGGTTGCTGTTTGGCTCGGTGGTCATCTTCTCCTCTGCTACCGAGAGCGGGTGCTAGAG
ACACAGCTGAGTGCTGGGGCGAGCGCGGGCATCGCTCTGGGCATCGGGCTGCTCTGCGGGCTGGTGGCCATGCTA
GTGCGCAGCGTGGGCCTCTTCTGGTGGGGCTGCTGCTCGGCCTGCTGCTCGCAGCTGCTGCCCTGCTGGGCTCC
GCACCCTACTACCAGCCAGGCTCCGTGTGGGGTCCACTGGGGCTGTGCTGGGGGGCGGCCTGCTCTGTGCCCTG
CTCACTCTGCGCTGGCCCCGCCACTCACCACCCTGGCCACCGCCGTGACTGGTGCTGCGCTGATCGCCACTGCC
GCTGACTACTTCGCGGAGCTGCTACTGCTGGGGCGCTACGTGGTGGAGCGACTCCGGGCTGCTCCTGTGCCCCCA
CTCTGCTGGCGAAGCTGGGCCCTGCTGGCACTCTGGCCCCTGCTCAGCCTGATGGGCGTTCTGGTGCAGTGGAGG
GTGACAGCTGAGGGGGACTCCACACGGAAGTGGTCATCAGCCGGCAGCGCCGACGCGTGCAACTGATGCGGATT
CGGCAGCAGGAAGATCGCAAGGAGAAAAGGCGGAAAAGAGACCTCCTCGGGCTCCCCCTCAGAGGTCCCCGGGCT
CCTCCCAGGCCTGGGCCACCAGACCCTGCTTATCGGCGCAGGCCAGTGCCCATCAAACGCTTCAATGGAGACGTC
CTCTCCCCGAGCTATATCCAGAGCTTCCGAGACCGGCAGACCGGGAGCTCCCTGAGCTCCTTCATGGCCTCACCC
ACAGATGCGGACTATGAGTATGGGTCCCGGGGACCTCTGACAGCCTGCTCAGGCCCCCAGTGCGGGTATAGCCA
TATCTGTCTGTCTAGACTCTGCAGTCACCAGCTCTGCCAGCTCGAGGAGGCCTGCTAGGCTGCCACTCAGCCTCC
TGGCTTTGGCTGTCCCTCTCCCCAGCCTGGAGAGGGCTGGCCTGGTCACTAGAAGGGAGGATTGTCTCAGGCCGAG
TCTTGGCCTGAGAGGAAAGCCCCCTCCCAAGCTCCCAAGAGGCTCCTGAGGAACTCGGGGTGTGAACCCATTGG
GGTGTGCTCAGGGTTGTGAGTGTGTTGCCCGTGTGCTGTGTGTATGTGTGTGGGGGTGGGCAGGCTTGGAGGGG
ACGCTGGGACCCTTGCCCTTAGATTTCTGACTGGTAGGGTTTCTCCAGGCTCAGCCCCACCTCTTCACTCCCTGCC
AAGGTCCCATGGGCCACCTCCTGCATGTCTCCGCGGAGGGGCTACCTTCCCTCCCATCGCCCTGCCTCGCAGCCA
GACTCATCTAAGGGTTCTTGTCTTGTCTATGGGGCAAACCTGTAGCATCCCTCACCTGGTCCCTGGCCTCTGT
AAAGCCACCAGCCTGAGGGCAGTGGCAGGAGATGGGGGTGGGGGGTGTGCTGCTGCGGCTGGGTGGGAAGGGAG
TTGGGGAGGGGTTTAAATGCACGGTGCATGTCTGGTGTCTGTGTCATGCCAACCTAGACACCTCATGCTTCTGTCTC
CCCCACCCACTCTGTTTTACATCTTTTATAAATGTGCCAACTGTGTGGCCTCTGCCA

1042/6881
FIGURE 972

MPGTVATLRFQLLPPEPDDAFWGAPCEQPLERRYQALPALVCIMCCLEFVVYCFFGYRCFKAVLFLTGLLEFGSVV
IFLLCYRERVLETQLSAGASAGIALGIGLLCGLVAMLVRSVGLFLVGLLLGLLLAAAALLGSAPYYQPGSVWGPL
GLLLGGGLLCALLTLRWPRPLTTLATAVTGAALIATAADYFAELLLLGRYVVERLRAAPVPPLCWRSWALLALWP
LLSLMGVLVQWRVTAEGDSHTEVVISRQRRRVQLMRIQQEDRKEKRRKKRPPRAPLRGPRAPPRPGPPDPAYRR
RPVPIKRFNGDVLSPSYIQSFRDRQTGSSLSSFMASTADYEGSRGPLTACSGPPVRV

1043/6881
FIGURE 973

CGGAGCACATGCTGAGCGGAGCGGCTGGGGCTGCGCGGCGTGGCGGAGCAGCGCTCGCTCCCTCGCTCACTCGCT
CGCTCGCAGGGACACACGCAGGGGCTGACAGCTGTGCTGGTGTCTGATAAGGGAAGCCACAAGGAGACGATCGAGG
AGAGAGACAAGCGGCAGCAGAGGCAGCAGCGGCAGAGGCAGCACCAGGGCTGCGGAGCTGCTGGGAGTGGGAGTG
ACTCCCCACCTCGGGCCCCCACCCTGTCCCTGTCTCTTCCCGCTTGGCCTGAGTTTAGAAGAGCAGCCGCTGC
CACCCTGCCACTCGGGAGGGCACCAGGGCTGCTGGCTAGGGAGGGACAGGGCAGGAGGCTCTGGCCAGTCCCA
GCAGCCGGGGACAGATGCCGATCGAGATTGTGTGCAAAATCAAATTTGCTGAGGAGGATGCGAAACCCAAGGAGA
AGGAGGCAGGGGATGAGCAGAGCCTCCTCGGGGCTGTTGCCCTGGAGCAGCCCCCGAGACCTGGCCACCTTTG
CCAGCACCAGCACCCTGCATGGACTGGGCCGGGCTGTGGCCAGGCCCCACGGACTGCGCAGAACCCTGTGGG
CACTGGCCCTACTCACCTCGCTGGCTGCCTTCTGTACCAGGCGGCTGGCCTGGCCCGGGGCTACCTGACCCGGC
CTCACTGGTGGCAATGGACCCCGCTGCCCCAGCCCCAGTGGCGGGCTTCCCGGCTGTACCCCTCTGCAATATCA
ACCGCTTCCGGCATTCCGGCACTCAGCGATGCCGACATCTTCCACCTGGCCAATCTGACAGGGCTGCCCCCAAAG
ACCGGATGGGCACCGTGGCGCTGGCTGCGCTACCCAGAGCCTGACATGGTAGACATCTCAACCGCACTGGCC
ACCAGCTCGCCGACATGCTTAAGAGTGCAACTTCAGTGGGCATCACTGCTCCGCCAGCACTTCTCTGTGGTCT
ATACTCGCTATGGGAAGTGTTACACCTTCAACGCGGACCCGCGGAGCTCGCTGCCAGCCGGGCAGGGGGCATGG
GCAGTGGCCTGGAGATCATGCTGGACATCCAGCAGGAGGAGTACCTGCCCATCTGGAGGGAGACAAATGAGACGT
CGTTTGAGGCAGGTATTTCGGGTGCAGATCCACAGCCAGGAGGAGCCGCCCTACATCCACCAGCTGGGGTTTCGGG
TGCCCCAGGCTTCAGACCTTTGTGTCTGCCAGGAACAGCGGCTGACCTACCTGCCCCAGCCCTGGGGCACT
GCCGCGCAGAGAGTGAGCTCAGGGAGCCTGAGCTTCAGGGCTACTCGGCCTACAGTGTGTCTGCCTGCCGGCTGC
GCTGTGAAAAGGAGGCCGTGCTTCAGCGCTGCCACTGCCGGATGGTGCACATGCCAGGCAATGAGACCATCTGCC
CACCAAATATCTACATCGAGTGTGCAGACCACACACTGGACTCCCTGGGTGGGGGCCCTGAGGGCCCGTGTCTTCT
GCCCCACCCCTGCAACCTGACACGCTATGGGAAAGAGATCTCCATGGTCAGGATCCCCAACAGGGGCTCAGCCC
GGTACCTGGCGAGGAAGTACAACCGCAACGAGACCTACATACGGGAGAACTTCTTGGTCTTAGATGTCTTCTTTG
AGGCCCTGACCTCTGAAGCCATGGAGCAGCGAGCAGCCTATGGCCTGTGAGCCCTGCTGGGAGACCTCGGGGGAC
AGATGGGCCTGTTCAATTGGGGCCAGCATCCTCACGTTGCTGGAGATCCTCGACTACATCTATGAGGTGTCTGGG
ATCGACTGAAGCGGTATGGAGGCGTCCCAAGACCCCTGCGGACCTCCACTGGGGGCATCTCCACTTTGGGGC
TTCAGGAGCTGAAGGAACAGAGTCCCTGCCCGAGCCTGGGCCGAGCGGAGGGTGGGGGGGTCAGCAGTCTGCTCC
CCAATCACCACCACCCCCACGGTCCCCAGGAGGTCTCTTTGAAGATTTTGCTTGCTAGGACGGTGCTGTGACTG
AAAGGACCCAGGAGTCTGGGACCCCTCCTGGGATCCCCAGCACATTCTCCTGCTCCTGGGAGAGGCCTGGGGGCG
GTGCTCACTGGGAGGGCCAGGACTCAGTTCTGTCTCATCCTCCCTGCCCTGATGTCAGCTGCTTTGCACAAA
GGTCCTTCTTGTCCACACCCCTTATCCCCAGGCTGGTGGCCCGGAGGGCTGGAGACCAGGCCATGGGCCCTCAC
GGAGAGGAAGGAAGGAAGGAGAGGGAGGGGGAGGATAGAGCCCATCCAGCCGGGAGGGGGAGCCCTCTGTAC
ATTTGTAAATATTTAGGGAAAGCCGGGTGGGGGGAGGGGATACAGATGTAGAAGGTGGGTAGGGCTACAGGGGTG
GGTGATTTAGGGACAGCCAGGGTCCCAGCCCCAATGTGACGAGGATAGGGAGAGCCCAGGACTCAGGAGTGCTG
GGCTGGTCTACTTCTGCCCCCTCTCCAGGCCCAGCTCCCTCTTGGCAGGGGGAGAGGATGGCCAGCAGGCCCT
GGCCAGCTCCAGTTCCCCCTGCACCAGCCCCACCCCTAGAGTCCCTTCTATAGGGAGGGGGCAGGAGACCTTC
CAGACTTCGGCTGAGCTTGGAGGGTGGGAAGGGAGCCTTCTCAGTCTCTCTCCCTCCAGTCTGATTTTATAAAG
TGCTGACGAG

1044/6881
FIGURE 974

MPIEIVCKIKFAEEDAKPKEKEAGDEQSLLGAVAPGAAPRDLATFASTSTLHGLGRACGPGPHGLRRTLWALALL
TSLAFLYQAAGLARGYLTRPHLVAMDPAAPAPVAGFPVAVTLCNINRFRHSALSDADIFHLANLTGLPPKDRDGH
RAAGLRYPEPDMVDILNRTGHQLADMLKSCNFSGHHCSASNFSVVYTRYGKCYTFNADPRSSLPSRAGGMGSGLE
IMLDIQQEEYLPWRETNETSFEAGIRVQIHSQEEPPYIHQLGFGVSPGFQTFVSCQEQRITYLPQPWGNCRAES
ELREPELQGYSAISVSACRLRCEKEAVLQRCRCRMVHMPGNETICPPNIYIECADHTLDSLGGGPEGPCFCPTPC
NLTRYGKEISMVRIPNRGSARYLARKYNRNETYIRENFLVLDVFFEALTSEAMEQRAAYGLSALLGDLGGQMGLF
IGASILTLLEILDYIYEVSWDRLEKRVWRRPKTPLRTSTGGISTLGLQELKEQSPCPSLGRAEGGGVSSLLPNHHH
PHGPPGGFLFEDFAC

1045/6881
FIGURE 975

CCTCGCCGCATCCACTCTCCGGCCGGCCGCCTGCCCCGCCGCTCCTCCGTGCGCCCGCCAGCCTCGCCCGCGCCG
TCACCATGAGCCAGGCCTACTCGTCCAGCCAGCGCGTGTCTCTACCGCCGCACCTTCGGCGGGGCCCCGGGCT
TCCCACTCGGCTCCCCGCTGAGTTCGCCCCGTGTTCCCGCGGGCGGGTTTCGGCTCTAAGGGCTCCTCCAGCTCGG
TGACGTCCCGCGTGTACCAGGTGTGCGGCACGTCGGGCGGGGCGGGGGCCTGGGGTCGCTGCGGGCCAGCCGGC
TGGGGACCACCCGCACGCCCTCCTCTACGGCGCAGGCGAGCTGCTGGACTTCTCACTGGCCGACGCGGTGAACC
AGGAGTTTCTGACCACGCGCACCAACGAGAAGGTGGAGCTGCAGGAGCTCAATGACCGCTTCGCCAACTACATCG
AGAAGGTCCGCTTCTGGAGCAGCAGAACCGCGCGCTCGCCGCCGAAGTGAACCGGCTCAAGGGCCGCGAGCCGA
CGCGAGTGGCCGAGCTCTACGAGGAGGAGCTGCGGGAGCTGCGGCGCCAGGTGGAGGTGCTCACTAACCAGCGCG
CGCGCGTCGACGTGAGCGCGACAACCTGCTCGACGACCTGCAGCGGCTCAAGGCCAAGCTGCAGGAGGAGATT
AGTTGAAGGAAGAAGCAGAGAACAATTTGGCTGCGTTCCGAGCGGACGTGGATGCAGCTACTCTAGCTCGCATTG
ACCTGGAGCGCAGAATTGAATCTCTCAACGAGGAGATCGCGTTTCTTAAGAAAGTGCATGAAGAGGAGATCCGTG
AGTTGCAGGCTCAGCTTCAGGAACAGCAGGTCCAGGTGGAGATGGACATGTCTAAGCCAGACCTCACTGCCGCC
TCAGGGACATCCGGGCTCAGTATGAGACCATCGCGCTAAGAACATTTCTGAAGCTGAGGAGTGGTACAAGTCGA
AGGTGTCAGACCTGACCCAGGCAGCCAACAAGAACAACGACGCCCTGCGCCAGGCCAAGCAGGAGATGATGGAAT
ACCGACACCAGATCCAGTCTACACCTGCGAGATTGACGCCCTGAAGGGCACTAACGATTCCTGATGAGGCAGA
TGCGGGAATTGGAGGACCGATTTGCCAGTGAGGCCAGTGGCTACCAGGACAACATTGCGCGCCTGGAGGAGGAAA
TCCGGCACCTCAAGGATGAGATGGCCCCCATCTGCGCGAGTACCAGGACCTGCTCAACGTGAAGATGGCCCTGG
ATGTGGAGATTGCCACCTACCGGAAGCTGCTGGAGGGAGAGGAGAGCCGGATCAATCTCCCCATCCAGACCTACT
CTGCCCTCAACTCCGAGAAACCAGCCCTGAGCAAAGGGGTTCTGAGGTCCATACCAAGAAGACGGTGATGATCA
AGACCATCGAGACACGGGATGGGGAGGTGCTCAGTGAGGCCACACAGCAGCAGCATGAAGTGCTCTTAAAGACAGA
GACCTCTGCCACCAGAGACCGTCTCACCCCTGTCTCACTGCTCCCTGAAGCCAGCCTTCTTCCATCCCAGGA
CACCACACCCAGCCTCAGTCTCTCCCTCACAGCCTCTGACCCCTCCTCACTGGCCATCCCTCGTGGTCCCCAACA
GCGACATAGCCCATCCCTGCCTGGTCACAGGGCATGCCCCGGCCACCTCTGCGGACCCAGCTGTGAGCCTTGGC
TGTTGGCAGTGAGTGAGCCTGGCTCTTGCTGCTGGATGGAGCCCAGGCGGGAGCGGTGGCCCTGTCCCTCCCACCT
CTGTGACCTCAGGCACTAGCCTTTGGCTCTGGAGACAGCCCCAGAGCAGGGTGTGGGATACTGCAGGGCCAGGA
CTGAGCCCCGCGACCTCCCCAGCCCTAGCCCAGGAGAGAGAAAGCCAGGCAGGTAGCCAGGGGGACTAGCCCC
TGTGGAGACTGGGGGGCTTGAAATTGTCCCCGTGGTCTCTTACTTTTCTTTCCCAGCCCAGGGTGGACTTAGAA
AGCAGGGGCTACAAGAGGGAATCCCCGAAGGTGCTGGAGGTGGGAGCAGGAGATTGAGAAGGAGAGAAAGTGGGT
GAGATGCTGGAGAAGAGAGGAGAGGAGAGAGGAGAGCGGTCTCAGGCTGGTGGGAGGGGCGCCACCTCCCC
ACGCCCTCCCTCCCCTGCTGCAGGGGCTCTGGAGAGAAACAATAAAGAGATTACACACAAGCC

1046/6881
FIGURE 976

MSQAYSSSQRVSSYRRTFGGAPGFPLGSPLSSPVFPFRAGFGSKGSSSSVTSRVYQVSRTSGGAGGLGSLRASRLG
TTRTPSSYGAGELLDFSLADAVNQEFLLTTRTNEKVELQELNDRFANYIEKVRFLQQNAALAAEVNRLKGREPTR
VAELYEEELRELRRQVEVLTNQRARVDVERDNLLDDLQRLKAKLQEEIQLKEEAENNLAAFRADVDAATLARIDL
ERRIESLNEEIAFLKKVHEEEIRELQAQLQEQVQVEMDMSKPDLTAALRDIRAQYETIAAKNISEAEEWYKSKV
SDLTQAANKNNDALRQAKQEMMEYRHQIQSYTCEIDALKGINDSLMRQMRELEDRFASEASGYQDNIAARLEEEIR
HLKDEMARHLREYQDLLNVKMALDVEIATYRKLEGEESRINLPIQTYSALNFRETSPEQRGSEVHTKKTVMIKT
IETRDGEVVSEATQQQHEVL

1047/6881
FIGURE 977

GCACGAGGGTGATGAAGGCCTACGAGTGCGGCGCGGCCTGAAGGGGCACGCGGGGGACCTGCAAAGCTAGTGAGG
GGCGGGGCAGGCGGCGCGGTGGGGGCGGGCCGAGCCCCGAGGGCCAGATGAGCGGACACAGCCCCACGCGCGGGGC
CATGCAGGTGGCCATGAACGGTAAGGCCCGCAAAGAGGCGGTGCAGACTGCGGCTAAGGAACCTCTCAAGTTCGT
GAACCGGAGTCCCTCTCCTTTCCATGCTGTGGCTGAATGCCGCAACCGCCTTCTCCAGGCTGGCTTCAGTGAAC
CAAGGAGACTGAGAAATGGAATATTAAGCCCCGAGAGCAAGTACTTCATGACCAGGAACCTCTCCACCATCATAGC
TTTTGCTGTAGGGGGCCAGTACGTTCCCTGGCAATGGCTTCAGCCTCATCGGGGCCACACGGACAGCCCCGCTCCT
CCGGGTGAAACGTCGGTCTCGCCGAGCCAGGTGGGCTTCCAGCAAGTCGGTGTGGAGACCTATGGTGGTGGGAT
CTGGAGCACCTGGTTTGACCGTGACCTGACTCTGGCTGGACGCGTCATTGTCAAGTGCCCTACCTCAGGTCTGGCT
GGAGCAGCAGCTGGTGCACGTGGAGCGGCCATTCTTCGCATCCACACCTGGCCATCCATCTGCAGCGAAATAT
CAACGAGAACTTTGGGCCCCAACACAGAGATGCATCTAGTCCCCATTCTTGCCACAGCCATCCAGGAGGAGCTGGA
GAAGGGGACTCCTGAGCCAGGGCCTCTCAATGCTGTGGATGAGCGGCACCATTCGGTCTCATGTCCCTGCTCTG
TGCCCATCTGGGGCTGAGCCCCAAGGACATAGTGGAGATGGAGCTCTGCCTTGACAGACCCAGCCTGCGGTCTT
GGGTGGTGCCTATGATGAGTTCATCTTTGCTCCTCGGCTGGACAATCTGCACAGCTGCTTCTGTGCCCTGCAGGC
CTTGATAGATTCTGTGCAGGCCTGGCTCCCTGGCCACAGAGCCTCACGTGCGCATGGTCACACTCTATGACAA
CGAAGAGGTGGGTCTGAGAGTGCACAGGGAGCACAGTCACTGCTGACAGAGCTGGTGTGCGGCGGATCTCAGC
CTCGTGCCAGCACCCGACAGCCTTCGAGGAAGCCATACCCAAGTCTTCATGATCAGCGCAGACATGGCCCATGC
TGTGCATCCCAACTACCTGGACAAGCATGAGGAGAACCACGGCCTTTATTCCACAAGGGCCCCGTGATCAAGGT
GAACAGCAAGCAACGCTATGCTTCAAACGCGGTGTGAGAGGCCCTGATCCGAGAGGTGGCCAACAAAGTCAAGGT
CCCCCTGCAGGATCTCATGGTCCGGAATGACACCCCTGTGGAACCACCATTGGACCTATCTTGGCTTCTCGGT
GGGGCTGCGGGTGCTGGATTTAGGCAGCCCCAACTGGCCATGCACTCTATCCGGGAGATGGCCTGCACCACAGG
AGTCCTCCAGACCTCACCTCTTCAAGGGCTTCTTTGAGCTGTTCCCTTCTCTAAGCCATAATCTCTTAGTGGA
TTGAGCCCTCTTGAAAGACTTCTCTGCCATCCCTTTGCACCTGAGAGGGGAAGTTCTCAGCTGAGCTGAAGCTG
GATTATTAAAGTGGATTGTCACTCAGAAAAAAAAAAAAAAAAAAAAA

1048/6881
FIGURE 978

MQVAMNGKARKEAVQTAAKELLKFVNRSPSPFHAVAECRNRLQAGFSELKETEKWNIKPESKYFMTRNSSTIIA
FAVGGQYVPGNGFSLIGAHTDSPCLRVKRRSRRSQVGFQQVGVETYGGGIWSTWFDRLTLAGRVIVKCPTSGRL
EQQLVHVERPILRIPHLAIHLQRNINENFGPNTEMHLVPILATAIQEELEKGTPEPGPLNAVDERHHSVLMSSLLC
AHLGLSPKDIVEMELCLADTQPAVLGGAYDEFIFAPRLDNLHSCFCALQALIDSCAGPGSLATEPHVRMVTLYDN
EEVGSESAQGAQSLLTELVLRRISASCQHPTAFEEAIPKSFMISADMAHAVHPNYLDKHEENHRPLFHKGPVIKV
NSKQRYASNAVSEALIREVANKVKVPLQDLMVRNDTPCGTTIGPILASRLGLRVLDLGSPQLAMHSIREMACTTG
VLQTLTLFKGFFELFPSLSHNLLVD

1049/6881
FIGURE 979

CTTTGCATTGTTCTCATCCGCCTCCTTGCTCGCCGCAGCCGCCTCCGCCGCGCGCCTCCTCCGCCGCCGCGGAC
TCCGGCAGCTTTATCGCCAGAGTCCCTGAACTCTCGCTTCTTTTTTAATCCCCTGCATCGGATCACC GGCGTGCC
CCACCATGTCAGACGCAGCCGTAGACACCAGCTCCGAAATCACCACCAAGGACTTAAAGGAGAAGAAGGAAGTTG
TGAAGAGGCAGAAAATGGAAGAGACGCCCCCTGCTAACGGGAATGCTAATGAGGAAAATGGGGAGCAGGAGGCTG
ACAATGAGGTAGACGAAGAAGAGGAAGAAGGTGGGGAGGAAGAGGAGGAGGAAGAAGAAGGTGATGGTGAGGAAG
AGGATGGAGATGAAGATGAGGAAGCTGAGTCAGCTACGGGCAAGCGGGCAGCTGAAGATGATGAGGATGACGATG
TCGATACCAAGAAGCAGAAGACCGACGAGGATGACTAGACAGCAAAAAAGGAAAAGTTAACTAAAAAAGG
GCCGCCGTGACCTATTCACCTCCACTTCCCGTCTCAGAATCTAAACGTGGTCACCTTCGAGTAGAGAGGCCCGC
CCGCCACCGTGGGCAGTGCCACCCGCAGATGACACGCGCTCTCCACCACCAACCCAAACCATGAGAATTTGCA
ACAGGGGAGGAAAAAGAACCAAACTTCCAAGGCCCTGCTTTTTTCTTAAAAGTACTTTAAAAAGGAAATTTG
TTTGTATTTTTTATTACATTTTATATTTTGTACATATTGTTAGGGTCAGCCATTTTTTAATGATCTCGGATGAC
CAAACAGCCTTCGGAGCGTTCTCTGTCCTACTTCTGACTTTACTTGTGGTGTGACCATGTTTATTATAATCTCA
AAGGAGAAAAAACCTTGTAAGGCAAAATGACAACAGAAAAACAATCTTATTCGAGCATTCCAGTAAC
TTTTTGTGTATGTACTTAGCTGTACTATAAGTAGTTGGTTTGTATGAGATGGTTAAAAAGGCCAAAGATAAAG
GTTTCTTTTTTTTCTTTTTTGTCTATGAAGTTGCTGTTATTTTTTTTGGCCTGTTTGATGTATGTGTGAAAC
AATGTTGTCCAACAATAAACAGG

1050/6881
FIGURE 980A

GCGGCAAGCCCCGCCTCAGCCCCCTCTGGCAGGCTCCCGCCAGCGTCGCTGCGGCTCCGGCCCCGGGAGCGAGCGCC
CGGAGCTCGGAAAGATGCGGGCGCCCGGGGGCTGGGGGTCTCGGGGGATCCGGGGTCTCCGGCTGCTCCTCT
GCCTCCTGCTGCTGAGCAGCCGCCCGGGGGGCTGCAGCGCCGTTAGTGCCACGGCTGTCTATTTGACCGCAGGC
TCTGCTCTCACCTGGAAGTCTGTATTTCAGGATGGCTTGTTTGGGCAGTGCCAGGTGGGAGTGGGGCAGGCCCCGGC
CCCTTTTGCAAGTCACCTCCCCAGTTCTCCAACGCTTACAAGGTGTGCTCCGACAACCTCATGTCCCAAGGATTGT
CCTGGCAGCATGACCTCACCCAGTATGTGATCTCTCAGGAGATGGAGCGCATCCCCAGGCTTCGCCCCCAGAGC
CCCGTCCAAGGGACAGGTCTGGCTTGGCACCCAAGAGACCTGGTCTGCTGGAGAGCTGCTTTTACAGGACATCC
CCACTGGCTCCGCCCCCTGTGCCCAGCATCGGCTTCCACAACCACCAAGTGGGCAAAGGTGGAGCTGGGGCCAGCT
CCTCTCTGTCCCCCTCTGCAGGCTGAGCTGTCCCCTCTCTTGAGGACCTGTGCTGCCCCCACAGCCTCCCC
ACCTTCACTGAGTTACGAACCTGCCTTGCTGCAGCCCTACCTGTCCACCAGTTTGGCTCCCGTGATGGCTCCA
GGGTCTCAGAGGGCTCCCCAGGGATGGTCAGTGTGCGCCCCCTGCCAAGGCTGAAGCCCCCTGCCCTCTTCAGCA
GAACTGCCTCCAAGGGCATATTTGGGGACCACCTGGCCACTCTACGGGGACCTTCCAGGGCCTTACCTGCCC
AGCTTTTTTCAAGACTCTGGGCTGCTCTATCTGCCCCAGGAGTTGCCAGCACCCAGCAGGGCCAGGGTGCCAAGGC
TGCCAGAGCAAGGGAGCAGCAGCCGGGCAGAGGACTCCCCAGAGGGCTATGAGAAGGAAGGACTAGGGGATCGTG
GAGAGAAGCTGCTTCCCCAGCTGTGCAGCCAGATGCGGCTCTGCAGAGGCTGGCCGCTGTGCTGGCGGGCTATG
GGGTAGAGCTGCGTCAGCTGACCCCTGAGCAGCTCTCCACACTCTGACCCTGTGCTGAGCTACTGCCAAGGGTG
CAGGAAGAAATCCGGGAGGGGTGTAAATGTTGGAGCTGATATCAAGAAAACAATGGAGGGGGCCGGTGGAGGGCA
GAGACACAGCAGAGCTTCCAGCCCGCACATCCCCATGCCTGGACACCCCACTGCCAGCCCTACCTCCAGTGAAG
TCCAGCAGGTGCCAAGCCCTGTCTCCTCTGAGCCTCCCAAAGCTGCCAGACCCCTGTGACACCTGTCTGCTAG
AGAAGAAAAGCCCACTGGGCCAGAGCCAGCCCACGGTGGCAGGACAGCCCTCAGCCCGCCAGCAGCAGAGGAAT
ATGGCTACATCGTCACTGATCAGAAGCCCCCTGAGCCTGGCTGCAGGAGTGAAGCTGCTGGAGATCCTGGCTGAGC
ATGTGCACATGTCTCAGGACGCTTCATCAACATCAGTGTGGTGGGACCAGCCCTCACCTTCCGCATCCGGCACA
ATGAGCAGAACCTGTCTTTGGCTGATGTGACCCAACAAGCAGGGCTGGTGAAGTCTGAACTGGAAGCACAGACAG
GGCTCCAAATCTTGACAGACAGGAGTGGGACAGAGGGAGGAGGCAGCTGCAGTCTTCCCCAAACTGCGCACAGCA
CCTCACCCATGCGCTCAGTGTGCTCACTCTGGTGGCCCTGGCAGGTGTGGCTGGGCTGCTGGTGGCTCTGGCTG
TGGCTCTGTGTGTCGGCAGCATGCGCGGCAGCAAGACAAGGAGCGCCTGGCAGCCCTGGGGCCTGAGGGGGCCC
ATGGTGACACTACCTTTGAGTACCAGGACCTGTGCCGCCAGCACATGGCCACGAAGTCCTTGTTCAACCGGGCAG
AGGGTCCACCGGAGCCTTCACGGGTGAGCAGTGTGCTCTCCAGTTCAGCGACGCAGCCAGGCCAGCCCCAGCT
CCCACAGCAGCACCCCGTCTCGGTGCGAGGAGCCGGCCCAAGCCAACATGGACATCTCCACGGGACACATGATTC
TGGCATACATGGAGGATCACCTGCGGAACCGGGACCGCCTTGCCAAGGAGTGGCAGGCCCTCTGTGCCATACCAAG
CAGAGCCAAACACCTGTGCCACCGCGCAGGGGGAGGGCAACATCAAAAAGAACCGGCATCCTGACTTCTTGCCCT
ATGACCATGCCCCGATAAACTGAAGGTGGAGAGCAGCCCTTCTCGGAGCGATTACATCAACGCCAGCCCCATTA
TTGAGCATGACCCTCGGATGCCAGCCTACATAGCCACGCAGGGCCCGCTGTCCCATACCATCGCAGACTTCTGGC
AGATGGTGTGGGAGAGCGGCTGCACCGTCATCGTCATGCTGACCCCGCTGGTGGAGGATGGTGTCAAGCAGTGTG
ACCGCTACTGGCCAGATGAGGGTGCCTCCCTCTACCACGTATATGAGGTGAACCTGGTGTGCGAGCACATCTGGT
GCGAGGACTTTCTGGTGGGAGCTTCTACCTGAAGAAGTGCAGACCCAGGAGACGCGCACGCTCACGCAGTTCC
ACTTCCTCAGCTGGCCGGCAGAGGGCACACCGGCCTCCACGCGGCCCTGCTGGACTTCCGCAGGAAGGTGAACA
AGTGCTACCGGGGCCGCTCCTGCCCATCATCGTGCACTGCAGTGATGGTGCGGGGAGGACCGGCACCTACATCC
TCATCGACATGGTCCTGAACCGCATGGCAAAGGAGTGAAGGAGATTGACATCGCTGCCACCTTGGAGCATGTCC
GTGACCAGCGGCTGGCCTTGTCCGCTCTAAGGACCAGTTTGAATTTGCCCTGACAGCCGTGGCGGAGGAAGTGA
ATGCCATCCTCAAGGCCCTGCCCCAGTGAGACCTGGGGCCCCCTTGGCGGGCAGCCAGCCTCTGTCCCTCTTTG
CCTGTGTGAGCATCTCTGTGTACCCACTCCTCACTGCCCCACCAGCCACCTCTTGGCATGCTCAGCCCTTCTTA
GAAGAGTCAGGAAGGGAAAGCCAGAAGGGGCACGCTGCCAGCCTCGCATGCCAGAGCCTGGGGCATCCCAGAG
CCCAGGGCATCCCATGGGGGTGCTGCAGCCAGGAGGAGAGGAAAGGACATGGGTAGCAATTCTACCCAGAGCCTT
CTCCTGCCTACATTCCTTGGCTGGCTCTCCTGTAGCTCTCCTGGGGTCTGGGAGTCCCTGAACATCTGTGTG
TGTCCCCCTATGCTCCAGTATGGAAGAATGGGGTGGAGGGTCGCCACACCCGGCTCCCCCTGCTTCTCAGCCCCG
GGCCTGCCTCTGACTCACACTTGGGCGCTCTGCCCTCCCTGGCCTCACGCCCAGCCTCCTCCCACCACCTCCCA
CCATGCGCTGCTCAACCTCTCTCCTTCTGGCGCAAGAGAACATTTCTAGAAAAAATACTTTTGTACCAAGTGTGA

1051/6881
FIGURE 980B

ATAAAGTTAGTGTGTTGTCTGTGCAGCTGCAAAAAAAAAAAAAAAAAAAAA

1052/6881
FIGURE 981

MRRPRRPGGLGGSGGLRLLCLLLSSRPGGCSAVSAHGCLFDRRLCSHLEVCIQDGLFGQCQVGVGQARPLLQV
TSPVLQRLQGVLRLQMSQGLSWHDDLTYVISQEMERIPRLRPPEPRPRDRSGLAPKRPGPAGELLLQDIPTGSA
PAAQHRLFPQPPVGKGGAGASSLSPLQAE LLPPLLEHLLLPQPPHPSLSYEPALLQPYLFHQFGSRDGSRVSEG
SPGMVSVGPLPKAEAPALFSRTASKGIFGDHPGHSYGDLPGPSAQLFQDSGLLYLAQELPAPSRARVPRLPEQG
SSSRAEDSPEGYEKEGLGDRGEKPASPAVQPDAAQLRLAAVLAGYGVELRQLTPEQLSTLLTLLQLLPKGAGRNP
GGVVNVGADIKKTMEGPVEGRDTAELPARTSPMPGHPTASPTISSEVQQVPSPVSSEPPKAARPPVTPVLLEKSP
LGQSQPTVAGQPSARPAEEYGYIVTDQKPLSLAAGVKLEILAEHVMSSGSFINISVVGPAITFRIRHNEQNL
SLADVTQQAGLVKSELEAQTGLQILQTGVGQREEAAVLPQTAHSTSPMRSVLLTLVALAGVAGLLVALAVALCV
RQHARQQDKERLAALGPEGAGDITTFEYQDLCRQHMATKSLEFNRAEGPPEPSRVSSVSSQFSDAAQASPSHSST
PSWCEEPAQANMDISTGHMILAYMEDHLNRDRLAKEWQALCAYQAEPNTCATAQEGEKNRNRHPDFLPYDHAR
IKLKVESSPSRSDYINASPIIEHDPRMPAYIATQGPLSHTIADFWQMVWESGCTVIVMLTPLVEDGVKQCDRYWP
DEGASLYHVYEVNLVSEHIWCEDFLVRSFYLNKVTQETRLTQFHFLSWPAEGTPASTRPLLDFFRKVNKCYRG
RSCPIIVHCSDGAGRTGTIILIDMVLNRMAKGVKEIDIAATLEHVRDQRPGLVRSKDQFEFALTAVAEVNAILK
ALPQ

1053/6881
FIGURE 982

AGTTCTCACTGAGACCTGTCAACCCGACTCAACGTGAGACGCACCGCCCGGACTCACCAATGCGTGAATGCATCTC
AGTCCACGTGGGGCAGGCAGGTGTCCAGATGGGCAATGCCTGCTGGGAGCTCTATTGCTTGGAACATGGGATTCA
GCCTGATGGGCAGATGCCCAGTGACAAGACCATTGGTGGAGGGGACGACTCCTTCACCACCTTCTTCTGTGAAAC
TGGTGCTGGAAAACAGTACCCCGGGCAGTTTTTGTGGATCTGGAGCCTACGGTCATTGATGAGATCCGAAATGG
CCCATACCGACAGCTCTTCCACCCAGAGCAGCTCATCACTGGGAAAAGAGGATGCTGCCAACAACTATGCCCGTGG
TCACTATAACCATTTGGCAAGGAGATCATTGACCCAGTGCTGGATCGGATCCGCAAGCTGTCTGACCAGTGCACAGG
ACTTCAGGGCTTCTGGTGTTCACAGCTTTGGTGGGGGCACTGGCTCTGGCTTCACCTCACTCCTGATGGAGCG
GCTCTCTGTTGACTATGGCAAGAAATCCAAGCTGGAATTCTCCATCTACCCAGCCCCCAGGTGTCTACAGCCGT
GGTTCGAGCCCTACAACCTCTATCCTGACCACCCACACCACCCTGGAGCACTCAGACTGTGCCTTCATGGTGGACAA
CGAAGCAATCTATGACATCTGCCGCCGCAACCTAGACATCGAGCGCCCAACCTACACCAACCTCAATCGCCTCAT
TAGCCAAATTGTCTCCTCCATCACAGCTTCTCTGCGCTTTGACGGGGCCCTCAATGTGGACCTGACAGAGTTCCA
GACCAACCTGGTGCCCTACCCTCGCATCCACTTCCCCCTGGCCACCTATGCACCAGTCATCTCTGCAGAAAAGGC
ATACCACGAGCAGCTGTCCGTGGCAGAGATCACCAATGCCTGCTTTGAGCCTGCCAACCAGATGGTAAAGTGTGA
TCCCCGGCACGGCAAGTACATGGCCTGCTGCCTGCTGTACCGTGGAGATGTGGTGCCCAAGGATGTCAACGCTGC
CATTGCCGCCATCAAGACCAAGCGCAGCATTTCAGTTTGTGGACTGGTGCCCCACAGGCTTCAAGGTTGGTATCAA
CTACCAGCCTCCCACTGTGGTGCCTGGGGGTGACCTGGCCAAGGTGCAGCGTGCCGTGTGCATGCTGAGCAACAC
GACCGCCATCGCCGAGGCCTGGGCCCGCTGGACCACAAGTTTCGACCTGATGTATGCCAAGAGGGCGTTTGTGCA
CTGGTATGTGGGTGAGGGCATGGAGGAGGGTGAGTTCTCCGAGGCCCGTGAGGATATGGCTGCCCTGGAGAAGGA
TTATGAGGAGGTGGGCATCGACTCCTATGAGGACGAGGATGAGGGAGAAGAATAAAGCAGCTGCCTGGAGCCTAT
TCACTATGTTTATTGCAAAATCCTTTTCAAATAAACAGTTTCTTGACCGGTAAAAAAAAAAAAAAAAAAAAA
AAAAAAAAAAAAA

1054/6881
FIGURE 983

MRECISVHVGQAGVQMGNACWELYCLEHGIQPDGQMPSDKTIGGGDDSF T T F F C E T G A G K H V P R A V F V D L E P T V I
DEIRNGPYRQLFHPEQLITGKEDAANNYARGHYTIGKEIIDPVLDRIKLS D Q C T G L Q G F L V F H S F G G G T G S G F T
SLLMERLSVDYGKKSLEFSIYPAPQVSTAVVEPYNSILTTITLEHSDCAFMVDNEAIYDICRRNL D I E R P T Y T
NLNRLISQIVSSITASLRFDGALNVDLTFQTNLVPYPRIHFPLATYAPVISA EKAYHEQLSVAEITNACFEPAN
QMVKCDPRHGKYMACCLLYRGDVVPKDVNA A I A I K T K R S I Q F V D W C P T G F K V G I N Y Q P P T V V P G G D L A K V Q R A V
CMLSNTTAIAEAWARLDHKFDL MYAKRA F V H W Y V G E G M E E G E F S E A R E D M A A L E K D Y E E V G I D S Y E D E D E G E E

1055/6881
FIGURE 984

GGGCCTGCAGTTGGCAGGAGGGTCCCGGGCCAGAGCCAGCGGGCCGTGCTGAGACGGCGTACGTGCCCTGCGT
GAGTGCCTGGCGGGCGGCGCTGCGCTAGGGGAGTGGGCGGTGAGGCCTGGTCCACGTGCGTCCCTTCCCGGGACC
CCCGCAGCTTGGCGCCAGCGGCTACGTGAGCCAAGGCACCCGGATGTCCGCGCCCTCTCCGAGTGACCACTG
CGGCCTCCGGTCCCGCAGTGCCCGCAGCCTCGGCCGGCGTCCACGCATTGCCATGGTGACTGTGGGCAACTACTG
CGAGGCCGAAGGGCCCGTGGGTCCGGCCTGGATGCAGGATGGCCTGAGTCCCTGCTTCTTCTTACGCTCGTGCC
CTCGACGCGGATGGCTCTGGGGACTCTGGCCTTGGTGCTGGCTCTTCCCTGCAGACGCCGGGAGCGGCCCGCTGG
TGCTGATTGCTGTCTTGGGGGGCCGGCCCTCGCATCTCTCCCTACGTGCTGCAGTGCTTCTGGCCACACTTCA
GGCGGGCGTGCCCTGGCCGGCCTGGCTGGCCGGGTGGGCACTGCCCGGGGGGCCCCACTGCCAAGCTATCTACT
TCTGGCCTCCGTGCTGGAGAGTCTGGCCGGCGCCTGTGGCCTGTGGCTGCTTGTCTGGAGCGGAGCCAGGCACG
GCAGCGTCTGGCAATGGGCATCTGGATCAAGTTACGGCACAGCCCTGGTCTCCTGCTCCTCTGGACTGTGGCGTT
TGCAGCTGAGAACTTGGCCCTGGTGTCTTGAACAGCCACAGTGGTGGTGGGCAAGGGCAGACTTGGGCCAGCA
GGTTCAGTTTACCTGTGGGTGCTGCGGTATGTGGTCTCTGGAGGGCTGTTTGTCTGGGTCTCTGGGCCCTGG
ACTTCGTCCCGAGTCTATACATTGCAGTTTCATGAAGAGGACCAAGATGTGGAAGGAGCCAGGTTGCGTCAGC
AGCCCAACAGTCTACCTGGCGAGATTTTGGCAGGAAGCTCCGCCTCCTGAGTGGCTACCTGTGGCCTCGAGGGAG
TCCAGCTCTGCAGTGGTGGTGCTCATCTGCCTGGGGCTCATGGGTTTGAACGGGCACTCAATGTGTGGTGCC
TATATTCTATAGGAACATTGTGAACCTGCTGACTGAGAAGGCACCTTGAACCTCTCTGGCCTGGACTGTTACCAG
TTACGTCTTCCCTCAAGTTCCCTCCAGGGGGGTGGCACTGGCAGTACAGGCTTCGTGAGCAACCTGCGCACCTTCCCT
GTGGATCCGGGTGCAGCAGTTCACGTCTCGGCGGGTGGAGCTGCTCATCTTCTCCACCTGCACGAGCTCTCACT
GCGTGGCACCTGGGGCGCCGCACAGGGGAGGTGCTGCGGATCGCGGATCGGGGCACATCCAGTGTACAGGGCT
GCTCAGCTACCTGGTGTTCATATGTATCCCCACGCTGGCCGACATCATATTGGCATCATCTACTTCAGCATGTT
CTTCAACGCCTGGTTTGGCCTCATTGTGTCTCTGTGCATGAGTCTTTACCTACCCCTGACCATTGTGGTCACTGA
GTGGAGAACCAAGTTTTCGTGCTATGAACACACAGGAGAAGCTACCCGGGCACGAGCAGTGGACTCTCTGCT
AACTTCGAGACGGTGAAGTATTACAACGCCGAGAGTTACGAAGTGAACGCTATCGAGAGGCCATCAATA
TCAGGGTTTGGAGTGAAGTCCGAGCGCTTCACTGGTTTTACTAAATCAGACCCAGAACCTGGTGATTGGGCTCGG
GCTCCTCGCCGGCTCCCTGCTTTGCGCATACTTTGTCACTGAGCAGAAGCTACAGGTTGGGGACTATGTGCTCTT
TGGCACCTACATTATCCAGCTGTACATGCCCCCAATTGGTTTGGCACCTACTACAGGATGATCCAGACCAACTT
CATTGACATGGAGAACATGTTTGACTTGCTGAAAGAGGAGACAGAAGTGAAGGACCTTCCCTGGAGCAGGGCCCCCT
TCGCTTTCAGAAGGGCCGTATTGAGTTTGAAGACGTGCACCTCAGCTATGCCGATGGGCGGAGACTCTGCAGGA
CGTGTCTTTCAGTGTATGCCCTGGACAGACACTTGCCCTGGTGGGCCCATCTGGGGCAGGGAAGAGCACAATTTT
GCGCCTGCTGTTTCGCTTCTACGACATCAGCTCTGGCTGCATCCGAATAGATGGGCAGGACATTTTACAGGTGAC
CCAGGCCTCTCTCCGGTCTCACATTGGAGTTGTGCCCAAGACACTGTCTCTTTAATGACACCATCGCCGACAA
TATCCGTTACGGCCGTGTACAGCTGGGAATGATGAGGTGGAGGCTGCTGCTCAGGCTGCAGGCATCCATGATGC
CATTATGGCTTTCCTGAAGGTTACAGGACACAGGTGGGCGAGCGGGGACTGAAGCTGAGCGCGGGGAGAAGCA
GCGCGTCGCCATTGCCCCGACCATCCTCAAGGCTCCGGGCATCAITCTGCTGGATGAGGCAACGTACGCGCTGGA
TACATCTAATGAGAGGGCCATCCAGGCTTCTCTGGCCAAAGTCTGTGCCAACCGACCCACCATCGTAGTGGCACA
CAGGCTCTCAACTGTGGTCAATGCTGACCAGATCCTCGTCATCAAGGATGGCTGCATCGTGGAGAGGGGACGACA
CGAGGCTCTGTTGTCCCGAGGTGGGGTGTATGCTGACATGTGGCAGCTGCAGCAGGGACAGGAAGAAACCTCTGA
AGACACTAAGCCTCAGACCATGGAACGGTGACAAAAGTTTGGCCACTTCCCTCTCAAAGACTAACCCAGAAGGGA
ATAAGATGTGTCTCTTCCCTGGCTTATTTTCATCCTGGTCTTGGGGTATGGTGCTAGCTATGGTAAGGGAAGG
GACCTTTCGAAAAACATCTTTTGGGGAAATAAAAAATGTGGACTGTG

1056/6881
FIGURE 985

GTGCACTTCAGCTATGCCGATGGGTGAGGCCCTTCCTTTTGCTTCCTTTGCTTTTCCGTTTCATTGACACTGCCT
TCCTGCTTCTCAGTGCTCTGAATCCCTGCTTTTGGAAAAGAGCCTGGGCAGGGGAGGGGCAGGGCCTACTAGCAG
CTCTGGTGATGGAAGGTGCCTGCTGAGCAACCCACCTTTCCTTGCAAGGCGGGAGACTCTGCAGGACGTGTCTTT
CACTGTGATGCCTGGACAGACACTTGCCCTGGTGAGAGGAGACCCAGCCACTTGGCCCAGATGCCTCAAGCTTCC
CTCATTTCAGTGCCCATGGTAGCTTGTGACCCAGGCTGTGGAGTCAGAGAGCCCAGTCACGATAGGCAACCGGATT
GAATGGTGCAGCCTCCATGGACACTGGTGACCTCTTGGAGAGGGAACCTCAAAGCCAGTGGGCCTCGGCCGGCT
GCGGTGGCACATGCCTGTAATCCCAGCACTTTGGGAGGCTTAGGTGGCCAGATCACCTAAGGTGGGCCCCATCTGG
GGCAGGGAAGAGCACAAATTTGCGCCTGCTGTTTCGCTTCTACGACATCAGCTCTGGCTGCATCCGAATAGATGG
G

1057/6881
FIGURE 986

GCTGCCCCGGTATCCCCGTGTGGGGCAGGAAGTGGGCTTCCAGATTCCCAGTATCCCCGGTAGGGTCTGCTTCTG
CTCGCTCGGGAAGGGCGGTGGCCGTCCCATTACTCAACCAACCCTCAACGCACTGGTCCTCCAGTTCCCTCCCC
TCAGTGGGTAAACAAACACACACCAGCGCTTGACTCGACAGACTCGAAAACAACATCTACTCAGAAAAGTTGTTT
CTTTCCTATCTCCTTAAACTTTCCCACTTCCACATATCGGAGCTTCTTCCACCTCGGCCTACTCAAGCATGAGA
TCGGAGGCGGGAGGGCGGCGACTGGCGGCGCGATGGACCTGACCGGGCTCCTGCTGGACGAAGAAGGCACCTTCT
CCCTCGCCGGCTTCCAGGACTTCACGTTCTTCCAGGACACCAGAAGCTGAGTCCCCGGATCCGAAGGAGGCTCT
ACTATGGCTGGGACTGGGAAGCCGACTGTAGCCTGGAGGAGCTTCCAGCCCGGTGGCAGACATTGCTGTGAAC
TGCTCCAGAAGGCAGCCCCCAGCCCTATTGCGCGACTCCAGAAGAAATATGTAGCTCATGTGTCCCGGAGGCAT
GCATCTCCCCATGTGCTATGATGCTGGCTCTGGTGTACATTGAACGGCTCCGGCACCAGAAACCCAGACTACTTGC
AGCATGTGTCATCCTCTGACTTGTTCCTGATCTCCATGATGGTGGCCAGTAAGTACCTCTATGATGAAGGGGAGG
AGGAGGAGGTCTTCAACGACGAATGGGGAGCTGCTGGGGGTGTGGCCGTGCCCACTCTCAATGCCTTGAGAGGG
GCTTCCTGAGTGCCATGGATTGGCATCTCTACACTGACCCTCGGGAGATCTTTGAGGTGCTGAGCTGGTTGGAGA
GCTGTGTGGCTGAGCAGCAGGGACGGTGGCGAGGCTGGTACACCTACACAGACCTGTGTGTGCTGCTGGAGCAGC
CGACCTGGCAGTTGGCCCTGGGCTCCCTCTGCCAGCGGCTGGTAAAGCTGTCTTGCCTGTTAGCTGTGGCATATG
TGAGCAGTGTGGCCCTGGCTGTGGCATCGGTGGCCGTAATACATCAGTCTTTGGGGCTGTCTGCATCCCTACAC
CTGGGCCCGCTGACCTTGGACTGACCTCCCGTTGCCTCCTGGAGCCCTGCATACCTTCTGTGCCACAATGCCTGC
CGTCTCTCGTAATGTCTCCAGCTGCCTGGAAGGCAGCATGGGGCTGCGGTCACTCTGGGGCAGTCTTCTGGCCT
CACTGACTCCTCCACCATTGCCTCCCCCAGACCCCCCTGCCCCCTCCCACTCTTCTTCATAACTGCCACCTTTGCC
AGAAGCTCCAGAGAGACTCCCCAACCTGCCATGCCTGCCTCCACCCCAACCGTACAGTCCCCACTGCGCTGTCCA
GCCCCCTGGTACCATAACCTATGGCCTGGCTCCCCCTGGCCTTGAGAGCCCGGTGCTCCTTTCACTTCCTCAGCCTC
AGCAATGTTCCCTTTTCACTGTGATGGAGCTGGCTCGCCTCAAGTCTTTTCGTTTTCCAGGCTAGGTAGGGCTGT
GAGGAATGCATTAAGAGGGTTTGGGAGTTTCTGAGAACCTGGAGGAGCAAAGCTTGATTGAGATCCTGTCTGCCT
CGCTGGGTCTTTGGCAGGTCCCCGTCTCCTCCTGGGTGGGAGCTTATGGGGTGGTGGGGCAGAAGGACTGAAGGTA
ATTCACTCCTAGATCGCAGTGGCTGGCTGCTTGGCCAGGACAGTGATGCCGCCAGGGAGAGCTTCCGCTTGGTGA
CCAGGGACATGTCCCAGATGGACATAGAAGCCCCCTCTCTGCCTCCCTGGGATTTTTTAGACTTTTACTTTTGATT
TCCCTAGGATGGAAGAGTATAGGTGGGAGATAAGGGAAGTGGGGTGAGAGGAGAAAGGAAATGTTGGCATGGGCC
TGTGTGATGTCCCTGAGGCAGAAGAGC

1058/6881
FIGURE 987

MDLTGLLLDEEGTFSLAGFQDFTFLPGHQKLSARIRRRLYYGWDWEADCSLEELSSPVADIARELLQKAAPSPIR
RLQKKYVAHVSREACISPCAMMLALVYIERLRHRNP DYLQHVSSDLFLISMMVASKYLYDEGEEEEVFNDEWGA
AGGVAVPTLNALERGFLSAMDWHLYTDPREIFEVL SWLESCVAEQQGRWRGWYTYTDL CVLLEQPTWQLALGSLC
QRLVKLSCLLAVAYVSSVALAVASVAVIHQSLGLSCIPTPGPPDLGLTSRCLLEPCIPSVPOCLPSLANVSSCLE
GSMGLRSLWGSLLASLTPPPLPPPDPAPPTLLHNCHLCQKLQ RDSPTCHACLHPNRTVPTALSSPWYHTYGLAP
PWPWSPVLLSLPQPQCSLFSVMELARLKSFVFPG

1059/6881
FIGURE 988

GAAGATATGGCGGCGTCTGCGTCTGCAGCTGCAGGGGAGGAGGACTGGGTCCTTCCCTCTGAAGTTGAAGTATTG
GAGTCCATCTATCTAGATGAACTACAGGTGATTAAAGGAAATGGCAGAACTTCACCATGGGAGATCTACATCACT
TTGCATCCTGCCACTGCAGAGGACCAGGATTCACAGTATGTCTGCCTCACTCTGGTGCTTCAGGTCCCAGCAGAG
TATCCCCATGAGGTGCCACAGATCTCTATCCGAAATCCCCGAGGACTTTCAGATGAACAGATCCACACGATCTTA
CAGGTGCTGGGCCACGTGGCCAAGGCTGGGCTGGGCACTGCCATGCTGTATGAACTCATTGAGAAAGGGAAGGAA
ATTCTCACAGATAACAACATCCCTCATGGCCAGTGTGTCTATCTGCCTCTATGGTTTCCAGGAGAAGGAGGCCTTT
ACCAAAACACCCTGTTACCACTACTTCCACTGCCACTGCCTTGCTCGGTACATCCAGCACATGGAGCAAGAGCTG
AAGGCACAAGGACAGGAGCAGGAACAGGAACGGCAGCATGCTACAACCAACAGAAGGCAGTCGGTGTGCAGTGT
CCAGTGTGCAGAGAGCCCCCTCGTGTATGATCTTGCCCTCACTGAAAGCAGCCCCCTGAACCCCAACAGCCCATGGAG
CTGTACCAGCCCAGTGCAGAGAGCTTGCGCCAGCAAGAAGAACGCAAGCGGCTCTACCAGAGGCAGCAGGAGCGG
GGGGGAATCATTGACCTTGAGGCTGAGCGAAACCGATACTTCATCAGCCTTCAGCAGCCTCCTGCCCCCTGCGGAA
CCTGAGTCAGCTGTAGATGTCTCAAAGGATCCCAACCAACCCAGCACCCCTTGCAAGCAGAACTATCCACCTCACCA
GCCGTCCAATCCACTTTGCCACCTCCTCTGCCTGTGGCGACCCAGCACATATGTGAGAAGATTCCAGGGACCAGG
TCAAATCAGCAAAGGTTGGGCGAAACCCAGAAAGCTATGCTAGATCCCCCAAGCCAGTCGAGGTCCCTGGCGA
CAGCCCGAACGGAGGCACCCAAAGGGAGGGGAGTGCCACGCCCCCTAAAGGTACCCGTGACACCCAGGAACCTGCCA
CCTCCTGAGGGGGCCCCCTCAAGGAGCCCATGGACCTAAAGCCAGAACCCCATAGCCAAGGAGTTGAAGGTCCCTCCA
CAAGAGAAGGGGCTTGGCAGCTGGCAGGGGGCCCCACCCCGCAGGACTCGGGACTGTGTTTCGTGGGAGCGCTCT
AAAGGCCGGACACCCGTTCTTCTTACCCTCGCCTGCCCTCGGGGCCAGTAGCATACCGGCTGGTACTCGGAGGG
AGTCCCTGGGCCTGGAATCTAAGGATGGTTCTTAGCAGGACTTGGTGGGGGAACAGGGAATTGGGGATGGGAGG
GAGGCAATAAAGATATTTGGCCTTC

1060/6881
FIGURE 989

MAASASAAAGEEDWVLPSEVEVLESIYLDELQVIKNGRTSPWEIYITLHPATAEDQDSQYVCFTLVLQVPAEYP
HEVPQISIRNPRGLSDEQIHTILQVLGHVAKAGLTAMLYELIEKGKEILTDNNIPHGQCVICLYGFQEKEAFTK
TPCYHYFHCHCLARYIQHMEQELKAQGQEQEQERQHATTKQKAVGVQCPVCREPLVYDLASLKAAPQPMPMELY
QPSAESLRQQEERKRLYQRQQERGGIIDLEAERNRYFISLQPPAPAEPESAVDVSKGSQPPSTLAAELSTSPAV
QSTLPPPLPVATQHICEKIPGTRSNQQRLEGETQKAMLDPPKPSRGPWRQPERRHPKGGECHAPKGTRDTQELPPP
EGPLKEPMDLKPEPHSQGVEGPPQEKGPSWQGPPPRRTRDCVRWERSKGRTPGSSYPRLPRGQ

1061/6881
FIGURE 990

GGCACGAGGGCCAGCTGCTGTAGAAAGAGGGGAGGAAACAAGCCAGTGCAAGGGGAGCAAAAGAGAAAAGGAGCCA
GGCTGGGCTTCTGATCCACAGCATCGCAGAGCTCGGGAGGCACAGCTCACAGACACAGGAAACACAGGACTGC
TATTCTGCTCTCCTGCCACGGTGATCTGGTGCCAGCTGGTGGAACAGTGGGTGATGGCGTCCCTGCTGCAAGAC
CAGCTGACCACTGATCAGGACTTGCTGCTGATGCAGGAAGGCATGCCGATGCGCAAGGTGAGGTCCAAAAGCTGG
AAGAAGCTAAGATACTTCAGACTTCAGAATGACGGCATGACAGTCTGGCATGCACGGCAGGCCAGGGGCAGTGCC
AAGCCCAGCTTCTCAATCTCTGATGTGGAGACAATACGTAATGGCCATGATTCCGAGTTGCTGCGTAGCCTGGCA
GAGGAGCTCCCCCTGGAGCAGGGCTTCACCAATTGTCTTCCATGGCCGCCGCTCCAACCTGGACCTGATGGCCAAC
AGTGTGAGGAGGCCAGATATGGATGCGAGGGCTCCAGCTGTTGGTGGATCTTGTCACCAGCATGGACCATCAG
GAGCGCTGGACCAATGGCTGAGCGATTGGTTTCAACGTGGAGACAAAATCAGGATGGTAAGATGAGTTTCCAA
GAAGTTCAGCGGTTATTGCACCTAATGAATGTGGAAATGGACCAAGAATATGCCTTCAGTCTTTTTCAGGCAGCA
GACACGTCCCAGTCTGGAACCCCTGGAAGGAGAAGAATTCGTACAGTTCTATAAGGCATTGACTAAACGTGCTGAG
GTGCAGGAAGTGTGAAAGTTTTTCAGCTGATGGGCAGAAAGCTGACTCTGCTGGAATTTTTGGATTTCCTCCAA
GAGGAGCAGAAGGAGAGAGACTGCACCTCTGAGCTTGCTCTGGAACCTATTGACCGCTATGAACCTTCAGACAGT
GGCAAACCTGCGGCATGTCTGAGTATGGATGGCTTCCTCAGCTACCTCTGCTCTAAGGATGGAGACATCTTCAAC
CCAGCTGCGCTCCCCATCTATCAGGATATGACTCAACCCCTGAACCACTACTTCATCTGCTCTTCTCATAACACC
TACCTAGTGGGGGACCAGCTTTGCGGCCAGAGCAGCGTCGAGGGATATATACGGGCCCTGAAGCGGGGTGCCGC
TGCGTGAGGTTGATGTATGGGATGGACCTAGCGGGGAACCTGTGCTTTACCACGGACACACCCCTGACCTCCCGC
ATCCTGTTCAAAGATGTCTGGCCACAGTAGCACAGTATGCCCTCCAGACATCAGACTACCCAGTCATCTTGTCC
CTGGAGACCCACTGCAGCTGGGAGCAGCAGCAGACCATGGCCCGTCATCTGACTGAGATCCTGGGGGAGCAGCTG
CTGAGCACCACCTTGGATGGGGTGTGCCCCACTCAGCTGCCCTCGCCTGAGGAGCTTCGGAGGAAGATCCTGGTG
AAGGGGAAGAAGTTAACTTGGAGGAAGACCTGGAATATGAGGAAGAGGAAGCAGAACCTGAGTTGGAAGAGTCA
GAATTGGCGCTGGAGTCCAGTTTGGAGCTGAGCCTGAGCCCCAGGAGCAGAACCTTCAGAATAAGGACAAAAAG
AAGAAATCCAAGCCCATCTTGTGTCCAGCCCTCTCTTCCCTGGTTATCTACTTGAAGTCTGTCTCATTCCGCAGC
TTCACACATTCAAAGGAGCACTACCACTTCTACGAGATATCATCTTTCTCTGAAACCAAGGCCAAGCGCCTCATC
AAGGAGGCTGGCAATGAGTTTGTGCAGCACATACTTGGCAGTTAAGCCGTGTGTATCCCAGCGGCTGAGGACA
GACTCTTCCAACATAACCCCCAGGAACCTGGAATGCAGGCTGCCAGATGGTGGCCATGAATATGCAGACTGCA
GGGCTTGAAATGGACATCTGTGATGGGCATTTCCGCCAGAATGGCGGCTGTGGCTATGTGCTGAAGCCAGACTTC
CTGCGTGATATCCAGAGTTCTTTCCACCCTGAGAAGCCCATCAGCCCTTTCAAAGCCCAGACTCTCTTAATCCAG
GTGATCAGCGGTGAGCAACTCCCCAAAGTGGACAAGACCAAGAGGGGTCCATTGTGGATCCACTGGTGAAAGTG
CAGATCTTTGGCGTTCTGTCTAGACACAGCACGGCAGGAGACCAACTATGTGGAGAACAATGGTTTTAATCCATAC
TGGGGGCAGACACTATGTTTCCGGGTGCTGGTGCTGAACTTGCCATGCTGCGTTTTGTGGTAATGGATTATGAC
TGGAATCCCGAAATGACTTTATTGGTCAGTACACCCTGCCCTGGACCTGCATGCAACAAGGTTACCGCCACATT
CACCTGCTGTCCAAAGATGGCATCAGCCTCCGCCAGCTTCCATCTTTGTGTATATCTGCATCCAGGAAGGCCTG
GAGGGGGATGAGTCTTGAGGTGGGCATTTACGGGAAGGGTTGGTATGCTGGCTTTAGACGGGGAGAAACATCTG
GAAGGATGCTCGAGAGAACAAATGGAGGTGGTGAAAATCAAGCTTTGGATTGTGCATTCCTAGGCACAAAATTAC
CTCATTCTTCTTAACAAGCAATCTGGGACCTGATTTCCACCTTTTTCTCTTTTCTTCCCTTCCCTTGTGTTTCA
TAAGCCTTTGGTATCTTTCTGCCCCTTTTCTTGTGTACTCTATACTGGAGTTCCCTTCTTCTCTTGTGCTGAG
GCTCAATCCCATACCGACATCTACAATAATCTTCCCATCAACTCTGTGTGAAGGCAGGTTGCAACTAGAAATT
CAGAGGGGCTTGGAATAGAGAAACCTAAAGAAGCATCATCCCTCCATCCCCAACTTCTCAAAGCCCAAAGCCA
AGGGAAGGATAAATCAAGGCTCAAGGCTTCCCCAGCAAAGATTAGGGAAAGAGACTTGACCCAGGACTGTACTA
CGACTCTTAAGAGAACTGCACAGCACTCAAAGTCCCCACTGGACTGCTTCTCCTTAGCCCCACTGGTATAA
ATACATCTCTCTCAATTTGGCAAAAAAAAAAAAAAAAAAAAA

1062/6881
FIGURE 991

MASLLQDQLTTDQDLLLMQEGMPMRKVRSKSWKKLRYFRLQNDGMTVWHARQARGSAKPSFSISDVETIRNGHDS
ELLRS�AEELPLEQGFTIVFHGRRSNLDLMANSVEEAQIWMRGLQLLDLVTSMDHQERLDQWLSDFQRGDKNQ
DGKMSFQEVQRLHLMNVEMDQEYAFSLFQAADTSQSGTLEGEFVQFYKALTKEAEVQELFESFSADGQKLTL
EFLDFLQEEQKERDCTSELALELIDRYEPSDSGKLRHVLSMDGFLSYLCSKDGDIENPACLPYQDMTQPLNHYF
ICSSHNTYLVGDQLCGQSSVEGYIRALKRGCRCEVDVWDGPGSEPVVYHGHTLTSRILEFKDVVATVAQYAFQTS
DYPVILSLETHCSWEQQQTMARHLTEILGEQLLSTLDGVLPTQLPSPEELRRKILVKGKLTLEEDLEYEEEEA
EPELESELALESQFETEPEPQEQNLQNKDKKKKSKPILCPALSSLYLKSVSFRSFTHSKEHYHFYEISSFSE
TKAKRLIKEAGNEFVQHNTWQLSRVYPSGLRTDSSNYPQELWNAGCQMVAMNMQTAGLEMDICDGHFRQNGCG
YVLKPDFLRDIQSSFHPEKPISPFKAQTLLIQVISGQQLPKVDKTKESIVDPLVKVQIFGVRLDTARQETNYVE
NNGFNYPWGTLCFRVLVPELAMLRFFVMDYDWKSRNDFIGQYTLPWTCMQQGYRHIHLLSKDGISLRPASIFVY
ICIQEGLEGDES

1063/6881
FIGURE 992

TTCTCCCCAGGCTCACTCACCATGACCAAGCTGAGCGCCCAAGTCAAAGGCTCTCTCAACATCACCACCCCGG
GCTGCAGATATGGAGGATCGAGGCCATGCAGATGGTGCCTGTTCTTCCAGCACCTTTGGAAGCTTCTTCGATGG
TGA CTGCTACATCATCTGGCTATCCACAAGACAGCCAGCAGCCTGTCTATGACATCCACTACTGGATTGGCCA
GGACTCATCCCTGGATGAGCAGGGGGCAGCTGCCATCTACACCACACAGATGGATGACTTCCTGAAGGGCCGGGC
TGTGCAGCACC CGAGGTCCAGGGCAACGAGAGCGAGGCCTTCCGAGGCTACTTCAAGCAAGGCCTTGTGATCCG
GAAAGGGGGCGTGGCTTCTGGCATGAAGCACGTGGAGACCAACTCCTATGACGTCCAGAGGCTGTGTCATGTCAA
GGGCAAGAGGAACGTGGTAGCTGGAGAGGTAGAGATGTCTTGAAGAGTTTCAACCGAGGGGATGTTTTCTCTCT
GGACCTTGGGAAGCTTATCATCCAGTGAATGGACCGAAAGCACCCGTATGGAGAGACTCAGGGGCATGACTCT
GGCCAAGGAGATCCGAGACCAGGAGCGGGGAGGGCGACCTATGTAGGCGTGGTGGACGGAGAGAATGAATTGGC
ATCCCCGAAGCTGATGGAGGTGATGAACCACGTGCTGGGCAAGCGCAGGGAGCTGAAGGCGCCGTGCCCCGACAC
GGTGGTGGAGCCGGCACTCAAGGCTGCACTCAAAGTGTACCATGTGTCTGACTCCGAGGGGAATCTGGTGGTGAG
GGAAGTCGCCACACGGCCACTGACACAGGACCTGCTCAGTCACGAGGACTGTTACATCCTGGACCAGGGGGCCCT
GAAGATCTACGTGTGAAAGGGAAGAAAGCCAATGAGCAGGAGAAGAAGGGAGCCATGAGCCATGCGCTGAACCT
CATCAAAGCCAAGCAGTACCCACCAAGCACACAGGTGGAGGTGCAGAAATGATGGGGCTGAGTCGGCCGTCTTTCA
GCAGCTCTTCCAGAAGTGGACAGCGTCCAACCGGACCTCAGGCCTAGGCAAAACCCACACTGTGGGCTCCGTGGC
CAAAGTGGAAACAGGTGAAGTTTCGATGCCACATCCATGCATGTCAAGCCTCAGGTGGCTGCCACGAGAAGATGGT
AGATGATGGGAGTGGGGAAGTGCAGGTGTGGCGCATTGAGAACCTAGAGCTGGTACCTGTGGATTCCAAGTGGCT
AGGCCACTTCTATGGGGCGGACTGCTACCTGCTGCTCTACACCTACCTCATCGGCGAGAAGCAGCATTACCTGCT
CTACGTTTGGCAGGGGACGCCAGGCCAGCCAAGATGAAATTACAGCATCAGCTTATCAAGCCGTCATCCTGGACCA
GAAGTACAATGGTGAACCAGTCCAGATCCGGGTCCCAATGGGCAAGGAGCCACCTCATCTTATGTCCATCTTCAA
GGGACGCATGGTGGTCTACAGGGGAGGCACCTCCCGAACTAACAACCTGGAGACCGGGCCCTCCACACGGCTGTT
CCAGGTCCAGGGAAC TGGCGCCAACAACACCAAGGCCTTTGAGGTCCAGCGCGGGCCAATTTCTCAATTCCAA
TGATGTCTTTGTCTCAAGACCCAGTCTTGCTGCTATCTATGGTGTGGGAAGGGTGTAGCGGGGACGAGCGGGA
GATGGCCAAGATGGTTGCTGACACCATCTCCCGACGGAGAAGCAAGTGGTGGTGGGAAGGGCAGGAGCCAGCCAA
CTTCTGGATGGCCCTGGGTGGGAAGGCCCCCTATGCCAACCAAGAGACTACAGGAAGAAAACCTGGTCATCAC
CCCCGGCTCTTTGAGTGTTCACAACAGACTGGGCGCTTCTGGCCACAGAGATCCCTGACTTCAATCAGGATGA
CTTGAAGAGGATGATGTGTTCTACTAGATGTCTGGGACCAGGTCTTCTTCTGGATTGGGAACATGCCAACGA
GGAGGAGAAGAAGGCCGACGCAACCACTGCACAGGAATACCTCAAGACCCATCCAGCGGGCGTGACCCTGAGAC
CCCCATCATTGTGGTGAAGCAGGGACACGAGCCCCCACCTTCACAGGCTGGTTCCTGGCTTGGGATCCCTTCAA
GTGGAGTAACACCAATCCTATGAGGACCTGAAGGCGGAGCTTGGCAACTCTAGGGACTGGAGCCAGATCACTGC
TGAGGTCAACAAGCCCCAAAGTGGACGTGTTCAATGCTAACAGCAACCTCAGTTCTGGGCCTCTGCCCATCTTCCC
CCTGGAGCAGCTAGTGAACAAGCCTGTAGAGGAGCTCCCCGAGGGTGTGGACCCAGCAGGAAGGAGGAACACCT
GTCCATTGAAGATTTCACTCAGGCCTTTGGGATGACTCCAGCTGCCTTCTCTGCTCTGCCTCGATGGAAGCAACA
AAACCTCAAGAAAGAAAAGGACTATTTTGAGAAGAGTAGCTGTGGTTGTAAAGCAGTACCCTACCCTGATTGTA
GGGTCTCATTTTCTACCGATATTAGTCCTACACCAATTGAAGTGAATTTTGCAGATGTGCCTATGAGCACAAA
CTTCTGTGGCAAATGCCAGTTTGTTTAATAATGTACCTATTCTTCAGAAAGATGATACCCC

1064/6881
FIGURE 993

TGCTCCCTTTTTCTGTCTCTGTCCATGCTGCCATGTTTCTCTGCTGCCAAATTGGGCCCCTTGGCCCCTTCCGGT
TCTGCTTCCTGGGGGCAGGGTTCCCTGCCTTGGACCCCCAGTCTGGGAACGGTGGACATCAAGTGCCTTGCATAGA
GCCCCCTCTTCCCCGCCAGCTTTCCAGGGGCACAGCTCTAGGCTGGGAGGGGAGAACCAGCCCCCTCCCCCTGC
CCCACCTCCTCTCCCCAGGGTCTAACGGTTAAGGGGACCCACATACCAGTGCCAAGGGGGATGTCAAGTGGTGAT
GTCGTTGTGCTCCCCCTCCCCAGAGCGGGTGGGCGGGGGTGAATATGGTTGGCCTGCATCAGGTGGCCTTCCCA
TTTAAGTGCCTTCTCTGTGACTGAGAGCCCTAGTGTGATGAGAACTAAAGAGAAAGCCAGACCCCT

1065/6881
FIGURE 994

ATGGACTCTGGGAGGCGTTTGGGCCAGAGAAGTGGATCCGCCGCTTGCGCCGATGGAGTCCGAATCGGAAAGC
GGGGCTGCTGCTGACACCCCCCACTGGAGACCCTAAGCTTCCATGGTGATGAAGAGATTATCGAGGTGGTAGAA
CTTGATCCCGGTCCGCCGGACCCAGATGACCTGGCCCAGGAGATGGAAGATGTGGACTTTGAGGAAGAAGAGGAG
GAAGAGGGCAACGAAGAGGGCTGGGTTCTAGAACCCAGGAAGGGGTGGTCGGCAGCATGGAGGGCCCCGACGAT
AGCGAGGTACCTTTGCATTGCACTCAGCATCTGTGTTTTGTGTGAGCCTGGACCCCAAGACCAATACCTTGGA
GTGACCGGGGGTGAAGATGACAAAGCCTTCGTATGGCGGCTCAGCGATGGGGAGCTGCTCTTTGAGTGTGCAGGC
CATAAAGACTCTGTGACTTGTGCTGGTTTCAGCCATGACTCCACTCTAGTGGCCACAGGGGACATGAGTGGCCTC
TTGAAAGTGTGGCAGGTGGACACTAAGGAGGAGGTC⁺TGGTCCTTTGAAGCGGGAGACCTGGAGTGGATGGAGTGG
CATCCTCGGGCACCTGTCCTGTTGGCGGGCACAGCTGACGGCAACACCTGGATGTGGAAAGTCCCGAATGGTGAC
TGCAAGACCTTCCAGGGTCCCAACTGCCCAGCCACCTGTGGCCGAGTCCTCCCTGATGGGAAGAGAGCTGTGGTA
GGCTATGAAGATGGGACCATCAGGATTTGGGACCTGAAGCAGGGGAAGCCCTATCCATGTACTGAAAGGGACTGAG
GGTACCCAGGGCCCCACTCACCTGTGTTGCTGCCAACCAGGATGGCAGCTTGATCCTAACTGGCTCTGTGGACTGC
CAGGCCAAGCTGGTCAGTGCCACCACCGCAAGGTGGTGGGTGTTTTTAGACCTGAGACTGTGGCCTCCAGCCC
AGCCTGGGAGAAGGGGAGGAGAGTGAGTCCAACCTCGGTGGAGTCCTTGGGCTTCTGCAGTGTGATGCCCCCTGGCA
GCTGTTGGCTACCTGGATGGGACCTTGCCATCTATGACCTGGCTACGCAGACTCTTAGGCATCAGTGTGAGCAC
CAGTCGGGCATCGTGCAGCTGCTGTGGGAGGCAGGCACTGCCGTGGTATATACCTGCAGCCTGGATGGCATCGTG
CGCCTCTGGGACGCCCGGACCGGCCGCTGCTTACTGACTACCGGGGCCACACGGCTGAGATCCTGGACTTTGCC
CTCAGCAAAGATGCCTCCCTGGTGGTGACCACGTGAGGAGACCACAAAGCGAAAGTATTTTGTGTCCAAAGGCCT
GACCGTTAATGGCTGCAGCCCCCTGCCTGTGTGTCTGGTGTGAGGGGACGAAGGGACCCCTGCCCTGTCTGCCA
GCAGAGGCAGTAGGGCACAGAGGGAAGAGGAGGGTGGGGCCCTGGATGACTTTCCAGCCTCTTCAACTGACTTGC
TCCCTCTCCTTTTTCTTCTTTAGAGACCCAGCCCAGGGCCCTCCCACCCTTGCCCAGACCTGGTGGGCCCTTC
AGAGGGAGGGGTGGACCTGTTTCTTTCACTTTCAATTTGCTGGTGTGAGCCATGGGGTGTGTATTTGTATGTGG
GGAGTAGGTGTTTGAAGTTCCCGTTCTTTCCCTTCCCAAGTCTCTGGGGTGGAAAGGAGGAAGAGATACTAGTT
AAAGATTTTAAAAATGTAAATAAAATATACTTCCAG

1066/6881
FIGURE 995

MDSGRRLGPEKWIRRLRMESESESGAAADTPPLETLSFHGDEEIIIEVVELDPGPPDPDDLAQEMEDVDFEEEE
EEGNEEGWVLEPQEGVVGSMEGPDSEVTFALHSASVFCVSLDPKTNLAVTGGEDDKAFVWRLSDGELLFECAG
HKDSVTCAGFSHDSTLVATGDMGSLKLVQVDTKEEVWSFEAGDLEWMEWHPRAPVLLAGTADGNTWMWKVPNGD
CKTFQGPNCPATCGRVLPDGKRAVVGIEDGTIRIWDLKQGSP IHVLKGTEGHQGP LTCVAANQDGS LILTGSVDC
QAKLVSATTGKVVG VFRPETVASQPSLGEGESESNVESLGFC SVMPLAAVGYLDGTLAIYDLATQTLRHQCQH
QSGIVQLLWEAGTAVVYTCSLDGIVRLWDARTGRLLTDYRGHTAEILDFALSKDASLVVTTSGDHKAKVFCVQRP
DR

1067/6881
FIGURE 996

GGCACGAGCTCTCCCTCCGTCCTTGCCCTCCCTTACCCACCTCACCGGCCCTTGTTTCTCCTTCCCCTGGGGGCA
GCCGCCGCCATGATCCTGCTGGAGGTGAACAACCGCATCATCGAGGAGACGCTCGCGCTCAAGTTCGAGAACGCG
GCCGCCGGAACAAACCGGAAGCAGTAGAAGTAACATTTGCAGATTTTCGATGGGGTCCTCTATCATATTTCAAAT
CCTAATGGAGACAAAACAAAAGTGATGGTCAGTATTTCTTTGAAATTCTACAAGGAAC TTCAGGCACATGGTGCT
GATGAGTTATTAAGAGGGTGTACGGGAGTTTCTTGGTAAATCCAGAATCAGGATACAATGTCTCTTTGCTATAT
GACCTTGAAAATCTTCCGGCATCCAAGGATTCATTGTGCATCAAGCTGGCATGTTGAAGCGAAATTGTTTTGCC
TCTGTCTTTGAAAATACTTCCAATTCCAAGAAGAGGGCAAGGAAGGAGAGAACAGGGCAGTTATCCATTATAGG
GATGATGAGACCATGTATGTTGAGTCTAAAAAGGACAGAGTCACAGTAGTCTTCAGCACAGTGTTTAAGGATGAC
GACGATGTGGTCATTGAAAAGGTGTTTCATGCAGGAGTTCAAAGAAGGACGCAGAGCCAGCCACACAGCCCCACAG
GTCCTCTTTAGCCACAGGGAACCTCCTCTGGAGCTGAAAGACACAGACGCCGCTGTGGGTGACAACATTGGCTAC
ATTACCTTTGTGCTGTTCCCTCGTCACACCAATGCCAGTGCTCGAGACAACACCATCAACCTGATCCACACGTTT
CGGGACTACCTGCACTACCACATCAAGTGCTCTAAGGCCTATATTCACACACGTATGCGGGCGAAAACGTCTGAC
TTCCTCAAGGTGCTGAACCGCGCACGCCCAGATGCCGAGAAAAAGAAATGAAAACAATCACGGGGAAGACGTTT
TCATCCCGCTTAATCTTGGGAATAAGAGGAGGAAGCGGCTGGCAACTGAAGGCTGGAACACTTGCTACTGGATAAT
CGTAGCTTTTAATGTTGCGCCTCTTCAGGTTCTTAAGGGATTCTCCGTTTTGGTTCCATTTTGTACACGTTTGGA
AAATAATCTGCAGAAACGAGCTGTGCTTGCAAAGACTTCATAGTTCCCAAGAATTAAAAAAAAAAAAAAAAAAAA

1068/6881
FIGURE 997

MILLEVNNRIIEETLALKFENAAAGNKPEAVEVTFADFDGVLYHISNPNGDKTKVMVSI SLKFYKELQAHGADEL
LKR VYGSFLVNPESGYNVSLLYDLENLPASKDSIVHQAGMLKRNCFASVFEKYFQFQEEGKEGENRAVIHYRDE
TMYVESKKDRVTIVVFSTVFKDDDDVVIGKVF MQEFKEGRRASHTAPQVLF SHREPPLELKDTDAAVGDNIGYITF
VLFPRHTNASARDNTINLIHTFRDYLHYHIKCSKAYIHTMR AKTSDFLKVLNRARPDAEKKEMKTI TGKTFSSR

1069/6881
FIGURE 998

GCGGCGGCGAGAGGATGAACAACAAGTTCGACGCTTTGAAAGATGATGACAGTGGGGACCATGATCAGAATGAAG
AAAACAGCACACAGAAAGATGGTGAGAAGGAAAAACGGAACGAGACAAGAATCAGAGCAGTAGCAAGAGAAAAGG
CTGTTGTCCCTGGACCGGCAGAGCATCCCCTGCAGTACAACCTACACTTTTGGTACTCCAGGAGAACCCCGGCC
GTCCCACGAGCTCACAGAGCTATGAACAGAATATCAAACAGATTGGCACCTTGCCTCTGTGGAGCAGTTCTGGA
GGTTTTATAGCCACATGGTACGTCCTGGGGACCTGACAGGCCACAGTGACTTCCATCTCTTCAAAGAAGGAATTA
AACCCATGTGGGAGGATGATGCAAAATAAAATGGTGGCAAGTGGATTATTCGGCTGCGGAAGGGCTTGGCCTCCC
GTTGCTGGGAGAACTCATTTTGGCCATGCTGGGGGAACAGTTCATGGTTGGGGAGGAGATCTGTGGGGCTGTGG
TGTCTGTCCGCTTTCAGGAAGACATTATTTCAATATGGAATAAGACTGCCAGTGACCAAGCAACCACAGCCCGAA
TCCGGGACACACTTCGGCGAGTGCTTAACCTACCTCCCAACACCATTATGGAATACAAAACCTCACACCGACAGCA
TCAAAATGCCAGGCAGGCTGGGCCCCCAAAGGCTCCTTTTCAAACCTCTGGAAGCCGCGGTTGAATGTGCCAT
GACCCCTCTCCCTCTCTGGATGGCACCATCATTGAAGCTGGCGTCATCGGAGTCTGTTGTTCTGTTGGCGTGCTAC
CTGGAAGATCCTTCTGTGGTGGACAAGAGGAATTGGAAGAGCATTTTATGTTTTAAGAACAGGCTGACACGCAGC
AGCTACAACAACAGCTGAGATCACTTAATAAATGGTGCTAAACTCAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA

1070/6881
FIGURE 999

MNNKFDALKDDDSGDHDQNEENSTQKDGEKEKTERDKNQSSSKRKAVVPGPAEHPLQYNYTFWYSRRTPGRPTSS
QSYEQNIKQIGTFASVEQFWRFYSHMVRPGDLTGHSDFHLFKEGIKPMWEDDANKNGGKWIIRLRKGLASRCWEN
LILAMLGEQFMVGEEICGAVVSVRFQEDIISIWNKTASDQATTARIRDTLRRVLNLPNTIMEYKTHTDSEIKMPG
RLGPQRLLFQNLWKPRLNVP

1071/6881
FIGURE 1000

GGCACGAGGAGAGCCTCGAGCCTGCGAGGAGCGCGCCGCCAGCTCCCTGCGTCCCGTCCCGCGTCCCCGCG
TTCCCGCGTCTCGGATCCGCGGCCATGGCCAGTGAGGAGCTGGCGTGCAAGCTGGAGCGCCGGCTGCGGCGCGA
GGAGGCCGAGGAGAGTGGCCCCAGCTGGCTCCCTCGGCGCCCCAGCCCCGGAGCCCCAAGCCCCGAGCCGAGCC
TCCCGCCCGTGCGCCACGGCCAGCGCCGACGCGGAGCTGAGCGCCCAGCTGAGCCGGCGGCTGGACATCAACGA
GGGCGCTGCGCGGCCCCGGCGCTGCAGGGTCTTCAACCCCTACACGGAGTTCCCGGAGTTCAGCCGCCGCTCAT
CAAGGACCTGGAGAGCATGTTCAAACCTGTATGACGCTGGCGGGATGGCTTCATCGACCTGATGGAGCTGAAGCT
GATGATGGAGAAGCTGGGGGCCCCCAGACCCACCTGGGCCTGAAGAGCATGATCAAGGAGGTGGATGAGGACTT
CGATGGCAAGCTCAGCTTCCGGGAGTTCCTGCTCATTTCCACAAGGCCGCGGCAGGGGAGCTGCAGGAGGACAG
TGGGCTGATGGCGCTGGCAAAGCTTTCTGAGATCGATGTGGCCCTGGAGGGTGTCAAAGGTGCCAAGAATTCTT
TGAAGCCAAGGTCCAAGCCTTGTCATCGGCCAGTAAGTTTGAAGCAGAGTTGAAAGCTGAGCAAGATGAGCGGAA
GCGGGAGGAGGAGGAGAGGCGGCTCCGCCAGGCAGCCTTCCAGAAACTCAAGGCCAACTTCAATACATAGTCTCTG
CTGACCTTGCCCTCTGCCACAGCTGTGCCTCACAGATGCCCCGAGAAGAGATGACTAGGCATCTTCATCACTGC
TGTCGGTCCCTCCCTGAGCCAGCATCTCCATCCACCACCCCGTGCCAGCTCCCGTGCCAGCCTTCATTCTCTCC
AGTGTCCAAGCCCCTCCAGGAGGGTCTGGGGTGGGCCAGATGCCTGCCACCTCTGTCTCCTGCTCTGCTCCT
CTGCCCTTCTTATAGCCAGAACTTGTATCTTCTCAGCAACCTTCACTTTGTCTTGTCCCTTTACCATTCCCCAT
CAAAGAGTAGTCTGCTATATCAATTTGTGTAGATATGTCTGTCTTTTGGGTCTCAGAGAAAATGCCATTTTC
TCGGAGAATTCTCTGCACTCTCTCTGCTTCACATTCAACTCCCTGTTCTCATCTTTGGTAGGATTCTGCCAGT
TGCTTTTGCATCTTCTGTTCTGGGTAATGGTGGGTCTTAATGGAGGCTGGGTGGACCACTGCCCGTCCACTCTT
CAACAGGAGGAACAGCATGCCACCACAGTAACACACATTAGAGAAAGGACAGAGGTCTGCTCCTTCTGCCACCT
TTCTCCTGGCCCCCTTAGCATTCCCCCAGTCCCTCCCTCTTACCTTGCTCCGTCTATGTCTTCCAGCTCAGCCT
TTTCCCCACTCTTAAATACTGTACTACTTCACTGTAAGAACGAAAGAATAGTTAGGATACCAATGAGTAAAAGG
TTCTGTCTCACTCTGACTCTGTGCAAATTGTATTACAGTAGACCGCTGACGTTCCCAAGTGACAGATCCAGGGCC
TTTCAAACATCCCCAAAGTCATGGCCATACTACCATTAGCCAGTTTCTAACATCTGTTTCAGGGTATCCAGCTG
TAGATGTTCTTATCCCCATACTTGTGAGTTATTGGGGTTGCTCACAATACTAGGGGTTTTGTGTATTTTA
ACAAATATATCCTAATGTATATTTATTCTTTTTGTAAGTGTCTTTACAATAAGAAATCATCTGCCTTTC
TAAAAAAAAAAAAAAAAAAAAA

1072/6881
FIGURE 1001

MASEELACKLERRLRREEAEESGPQLAPLGAPAPEPKPEPEPPARAPTASADAELSAQLSRRLDINEGAARPRRC
RVFNPYTEFPEFSRRLIKDLESMFKLYDAGRDGFIDLMELKLMMEKLGAPQTHLGLKSMIKEVDEDFDGKLSFRE
FLLIFHKAAAGELQEDSGLMALAKLSEIDVALEGVKGAKNFFEAKVQALSSASKFEAEKAEQDERKREEEERRL
RQAAFQKLKANFNT

1073/6881

FIGURE 1002

ATGTCCCCTGCTGGAACCATGGCAACATCACCCGCTCCAAGGCGGAGGAGCTGCTTTCCAGGACAGGCAAGGAC
GGGAGCTTCCTCGTGCGTGCCAGCGAGTCCATCTCCCGGGCATAACGCGCTCTGCGTGCTGTATCGGAATTGCGTT
TACACTTACAGAATTCTGCCCCAATGAAGATGATAAATTCAGTGTTCAGAAATCTGAACCTGACTTCAGCCGATAC
TGGCAGAGAAAGGGTGTAACGATGAGGAAGGAAGTGCACGCGCAGGATGCAGTAGTAGACAGGGAGCTGGACTC
AAGGTGTGGGGAGACCACAGGCAAGTCCCTGGAGGCAAAAATCCTCCAGAAGTTGTCAAGAGGGAGAGCGTGGTG
AGCAGAACGCCCCTCAGGAAGACAGCTTATGAGGATGGAAGGAATTGCTTCGAGGAGCGATGTGATTGA

1074/6881
FIGURE 1003

MVPCWNHGNITRSKAEELLSRTGKDGSLVRASESISRAYALCVLYRNCVYTYRILPNEDDKFTVQKSEPDFSRY
WQRKGVNDEEGSARAGCSSRQGAGLKVWGDHRQVPGGKNPPEVVKRESVVSRTPLRKTAYEDGRNCFEERCD

1075/6881
FIGURE 1004

CTTCGGGTGTACGTGCTCCGGGATCTTCAGCACCCGCGGCCGCCATCGCCGTCGCTTGGCTTCTTCTGGAATCAT
CTGCGCCACTTGTCCGCTTCACACTCCGCGGCCATCATGGTGAAGCTCGCGAAGGCAGGTAAAAATCAAGGTGAC
CCCAAGAAAATGGCTCCTCCTCCAAAGGAGGTAGAAGAAGATAGTGAAGATGAGGAAATGTCAGAAGATGAAGAA
GATGATAGCAGTGGAGAAGAGGTCGTCATACCTCAGAAGAAAGGCAAGAAGGCTGCTGCAACCTCAGCAAAGAAG
GTGGTCGTTTCCCCAACAAAAAGGTTGCAGTTGCCACACCAGCCAAGAAGCAGCTGTCACTCCAGGCAAAAAAG
GCAGCAGCAACACCTGCCAAGAAGACAGTTACACCAGCCAAAGCAGTTACCACACCTGGCAAGAAGGGAGCCACA
CCAGGCAAAGCATTGGTAGCAACTCCTGGTAAGAAGGGTGTGCCATCCCAGCCAAGGGGGCAAAGAATGGCAAG
AATGCCAAGAAGGAAGACAGTGATGAAGAGGAGGATGATGACAGTGAGGAGGATGAGGAGGATGACGAGGACGAG
GATGAGGATGAAGATGAAATTGAACCAGCAGCGATGAAAGCAGCAGCTGCTGCCCTGCCTCAGAGGATGAGGAC
GATGAGGATGACGAAGATGATGAGGATGACGATGACGATGAGGAAGATGACTCTGAAGAAGAAGCTATGGGAGACT
ACACCAGCCAAAGGAAAGAAAGCTGCAAAAGTTGTTCTGTGAAAGCCAAGAACGTGGCTGAGGATGAAGATGAA
GAAGAGGATGATGAGGACGAGGATGACGACGACGACGAAGATGATGAAGATGATGATGATGAAGATGATGAGGAG
GAGGAAGAAGAGGAGGAGGAAGAGCCTGTCAAAGAAGCACCTGGAAAACGAAAGAAGGAAATGGCCAAACAGAAA
GCAGCTCCTGAAGCCAAGAAACAGAAAGTGGAAAGGCACAGAACCGACTACGGCTTCAATCTCTTTGTTGGAAAC
CTAACTTTAACAATCTGCTCCTGAATTAATAAACTGGTATCAGCGATGTTTTTGTCTAAAAATGATCTTGCTGTT
GTGGATGTCAGAATTGGTATGACTAGGAAATTTGGTTATGTGGATTTTGAATCTGCTGAAGACCTGGAGAAAGCG
TTGGAACTCACTGGTTTGAAGTCTTTGGCAATGAAATTAAC TAGAGAAACCAAAGGAAAAGACAGTAAGAAA
GAGCGAGATGCGAGAACACTTTTGGCTAAAAATCTCCCTTACAAAGTCACTCAGGATGAATTGAAAGAAGTGTTT
GAAGATGCTCGGAGATCAGATTAGTCAGCAAGGATGGGAAAAGTAAAGGGATTGCTTATATTGAATTTAAGACA
GAAGCTGATGCAGAGAAAACCTTTGAAGAAAAGCAGGGAACAGAGATCGATGGGCGATCTATTTCCCTGTACTAT
ACTGGAGAGAAAGGTCAAAATCAAGACTATAGAGGTGGAAAGAATAGCACTTGGAGTGGTGAATCAAAAACCTCTG
GTTTTAAGCAACCTCTCCTACAGTGCAACAGAAGAACTCTTCAGGAAGTATTTGAGAAAGCAACTTTTATCAAA
GTACCCCAAGAACCAAAATGGCAAATCTAAAGGGTATGCATTTATAGAGTTTGCTTCATTGCAAGACGCTAAAGAA
GCTTTAAATTCTGTAAATAAAAGGGAAATTGAGGGCAGAGCAATCAGGCTGGAGTTGCAAGGACCCAGGGGATCA
CCTAATGCCAGAAGCCAGCCATCCAAAACCTCTGTTTGTCAAAGGCCTGTCTGAGGATACCACTGAAGAGACATTA
AAGGAGTCATTGACGGCTCCGTTCCGGGCAAGGATAGTTACTGACCGGGAAACTGGGTCTCCAAAGGGTTTGGT
TTTGTAGACTTCAACAGTGAGGAGGATGCCAAAGCTGCCAAGGAGGCCATGGAAGACGGTGAAATTGATGGAAAT
AAAGTTACCTTGGACTGGGCCAAACCTAAGGGTGAAGGTGGCTTCGGGGGTGCTGGTGGAGGCAGAGGCGGCTTT
GGAGGACGAGGTGGTGGTAGAGGAGGCCGAGGAGGATTTGGTGGCAGAGGCCGGGGAGGCTTTGGAGGGCGAGGA
GGCTTCCGAGGAGGCAGAGGAGGAGGAGGTGACCACAAGCCACAAGGAAAGAAGACGAAGTTTGAATAGCTTCTG
TCCCTCTGCTTTCCCTTTTCCATTTGAAAGAAAGGACTCTGGGGTTTTTACTGTTACCTGATCAATGACAGAGCC
TTCTGAGGACATTCCAAGACAGTATACAGTCCTGTGGTCTCCTTGGAAATCCGTCTAGTTAACATTTCAAGGGCA
ATACCGTGTTGGTTTTGACTGGATATTCATATAAACTTTTTAAAGAGTTGAGTGATAGAGCTAACCTTATCTGT
AAGTTTTGAATTTATATTGTTTCATCCCATGTACAAAACATTTTTCTCTAC

1076/6881
FIGURE 1005

MVKLAKAGKNQGDPPKMAPPPKEVEEDSEDEEMSEDEEDDSSGEEVVIPQKKGKKAATSARKVVSPTKKVAVA
TPAKKAAVTPGKKAAATPAKKTVTPAKAVTTPGKKGATPGKALVATPGKKGAAIPAKGAKNGKNAKKEDSDEEED
DDSEDEDEDEDEDEDEDEIEPAAMKAAAAAPASEDEDEDEDEDEDEDDDEEDDSEEEAMETTPAKGKKAAKV
PVKAKNVAEDEDEEEDDEDEDDDDDE
TEPTTAFNLFVGNLNFNKSAPELKTGISDVFAKNDLAVVDVRIGMTRKFGYVDFESAEDLEKALELTGLKVFGNE
IKLEKPKGKDSKKERDARTLLAKNLPYKVTQDELKEVFEDAAEIRLVSKDGKSKGIAYIEFKTEADAETFEKQ
GTEIDGRSISLYYTGEKGONQDYRGGKNSTWSGESKTLVLSNLSYSATEETLQEVFEKATFIKVPQNQNGKSKGY
AFIEFASFEDAKEALNSCNKREIEGRAIRLELQGPRGSPNARSQPSKTLFVKGLSEDTEETLKESFDGSSVRARI
VTDRETGSSKGFGFVDFNSEEDAKAAKEAMEDGEIDGNKVTLDWAKPKGEGGFGGGGGGGGFGGGGGGGGGGG
FGGRGRGGFGGGGGFRGGGGGGGDHKPQGKKTKE

1077/6881
FIGURE 1006

TGAACTGAGCGGCCCTGAGCTGACAGATACACTGCGCAGCTGGAACGGCGAGCGAGCCGACGGGCGAGTGAGGG
GCGCAGCCATGATCACCTCGGCCGCTGGAATTATTTCTCTTCTGGATGAAGATGAACCACAGCTTAAGGAATTTG
CACTACACAAATTGAATGCAGTTGTTAATGACTTCTGGGCAGAAATTTCCGAGTCCGTAGACAAAATAGAGGTTT
TATACGAAGATGAAGGTTTCCGGAGTCGGCAGTTTGCAGCCTTAGTGGCATCTAAAGTATTTATCACCTGGGGG
CTTTTGAGGAGTCTCTGAATTATGCTCTTGGAGCAGGGGACCTCTTCAATGTCAATGATAACTCTGAATATGTGG
AAACTATTATAGCAAAATGCATTGATCACTACACCAAACAATGTGTGAAAATGCAGATTTCCTGAAGGAGAAA
AAAAACCAATTGACCAGAGATTGGAAGGCATCGTAAATAAAATGTTCCAGCGATGTCTAGATGATCACAAGTATA
AACAGGCTATTGGCATTGCTCTGGAGACACGAAGACTGGACGTCTTTGAAAAGACCATACTGGAGTCAATGATG
TCCCAGGAATGTTAGCTTATAGCCTTAAGCTCTGCATGTCTTTAATGCAGAATAAACAGTTTCGGAATAAAGTAC
TAAGAGTTCTAGTTAAAATCTACATGAACCTTGGAGAAACCTGATTTCATCAATGTTTGTGAGTGTCTAATTTCT
TAGATGATCCTCAGGCTGTGAGTGATATCTTAGAGAACTGGTAAAGGAAGACAACCTCCTGATGGCATATCAGA
TTTGTGTTTGATTGTATGAAAGTGCTAGCCAGCAGTTTTTGTCATCTGTAATCCAGAATCTTCGAAGTGTGGCA
CCCCATTGCTTCTGTGCCTGGATCCACTAATACGGGTACTGTTCCGGGATCAGAGAAAGACAGTGACTCGATGG
AAACAGAAGAAAAGACAAGCAGTGCAATTTGTAGGAAAGACACCAGAAGCCAGTCCAGAGCCTAAGGACCAGACTT
TGAAAATGATTAAATTTTAAAGTGGTGAATGGCTATTGAGTTACATCTGCAGTTCTTAATACGAAACAATAATA
CAGACCTCATGATTCTAAAAAACACAAAGGATGCAGTACGGAATTCTGTATGTCTACTGCAACCGTTATAGCAA
ACTCTTTTATGCACTGTGGGACAACCAAGTGACCAAGTTTCTTAGAGATAATTGGAATGGTTAGCCAGAGCCACTA
ACTGGGCAAAATTTACTGCTACAGCCAGTTTGGGTGTAATTCATAAGGGTCATGAAAAAGAACGATTACAGTTAA
TGGCAACATACCTTCCCAAGGATACTTCTCCAGGATCAGCCTATCAGGAAGGTGGAGGTCTCTATGCACTAGGTC
TTATTTCATGCCAATCATGGTGGTGATATAATTGACTATCTGCTTAATCAGCTTAAGAAGCCAGCAATGATATCG
TTAGACACGGTGGCAGTCTGGGCCTTGTTTGGCAGCCATGGGAACGACGTCAAGATGTTTATGATTTGCTAA
AAACAAACCTTTATCAGGATGATGCAGTAACAGGGGAAGCAGCTGGCCTGGCCCTAGGTTTGGTTATGTTGGGCT
CTAAAAATGCTCAGGCTATTGAGGACATGGTTGGTTATGCACAAGAACTCAACATGAGAAGATTCTGCGTGGTC
TTGCAGTTGGCATAGCTTTAGTAATGTATGGGAGGATGGAAGAGGCTGATGCTCTCATTGAATCTCTCTGTCTGTG
ACAAGGACCCAATTTCTCGAAGGTCTGGAATGTATACGTAGCCATGGCTTATTGTGGCTCTGGTAACAACAAG
CAATTCGACGCCTGTACATGTTGCTGTAAGTGATGTTAATGATGATGTCAGGAGGGCAGCAGTAGAATCACTTG
GGTTCATTCTATTGAGAACCCTGAACAGTGCCCAAGTGTGTCTCTTTGTTGTGAGAGATTACAACCTCATG
TGCCTACGGAGCTGCAATGGCCTTGGGATATGCTGTGCTGGTACAGGAAACAAGGAAGCCATTAATTTGCTAG
AACCAATGACAAACGACCCCGTGAACACGTGAGGCAAGGGGCACTCATAGCTTCAGCTCTCATCATGATCCAGC
AGACTGAAATCACTTGTCCAAAGGTGAATCAGTTCAGACAGCTGTATTCCAAAGTCATCAATGATAAGCATGATG
ATGTCATGGCCAAGTTTGGCGCTATTCTGGCCAGGGCATACTGGATGCAGGTGGTCATAATGTCACAATCTCCT
TGCAGTCCAGGACTGGGCATACTCATATGCCTTCTGTGGTTGGCGTCTTGTATTTACCCAGTTTGGTTCTGGT
TTCTCTTTTACACTTCTGTCTATTGGCTTATACCCCTACCTGTGTCTATTGGCCTTAACAAGGACTTAAAGATGC
CGAAAGTTCAGTATAAATCGAACTGTAAACCATCCACATTTGCATATCCTGCCCTCTGGAAGTACCAAAAGAAA
AAGAAAAGGAAAAGGTTTCTACTGCTGTATTATCTATAACTGCCAAGGCTAAAAAGAGGAAAAAGAAAAGGAAA
AAAAGGAGGAGGAGAAAATGGAAGTGGATGAGGCAGAGAAAAAGGAGGAAAAAGAGAAGAAAAAGAACCTGAGC
CAAACCTCCAGTTATTGGATAACCCAGCCCGAGTTATGCCTGCCAGCTTAAGGTCCTAACCATGCCGAGACCT
GTAGATACCAGCCTTTCAAACCACTCTCTATTGGAGGCATCATCTTCTGAAGGATACCAAGTGAAGACATTGAGG
AGCTGGTGAACCTGTGGCAGCACATGGCCCAAAATCGAGGAGGAGGAACAAGAGCCAGAACCCCAAGAACCAT
TTGAGTATATTGATGATTAAGGGCCAGAGGATCTCACTTGCTTATCTGAAGAAGATTGTCCAGGCTCATATTGGG
AATGCTTATGAGGAAATTCATGCCGAGACCTGCTATTCAATGCATGTATCGTTGCCTCTGCACTGACCTGAAGAA
CCCTGTCTCCAAGTCTTTGGTTGAAGAGAAGATATGACTGTTGAGTGTGCTCTTTCACAGAACTTGGTTTTCA
AATAAATATAAGATCTCCAGATGGACAAG

1078/6881
FIGURE 1007

CTGCAAGCCGCCGGAGCCGGGCAACCAGTGGAAGGAGCCTGGGAGAGGCCAGGCCTCCCCGGACTGCTAGCCTGC
TTTTCTGGGGTCCCTGGAGCCGGAGGAAGAACCAGGATGTTGCTGCCTGCAGAAGCTCAGCTCAGGAAGACTTC
CAGGAACCTGAGGAGGAGCTGCCACTAACAGCCATATTTCCCAATGAGACTGTGATGACCTTGAAGGGGGTCA
AAAGCCTGTGATGGAGTCGTACACACTCCTGCTGAGCCCACCGGAGACTCAAGATGAAGGCTGGACCCTTGCCT
GTCCCTGGCTCTAACCTACAGACTGGGGCCTGGCTCCGTCTTACTGGCCCCAGGTCTCCATGGAGACTGCAGAA
ACCCCGCCTGCTGGAGGCCTGCCACACTACAGTTACCAGCTAGACAGTGGGGCTTACTAAGACAAGCAGGACC
TAAACAGTGTCTCCCTGGGAACCTACTCCCCACCCAGCATTTGCTAAGTCTGATCACAGGGAGGTTATTTTGT
CTCTGTGTCTCGGTTTCTCTGAGCCACTGAGACAGATGGCTGTCCGCTTTGAGGCTCTGCAGAGCTGTGGCACC
CATGGTGTGTCTGCAGTGTCTTGGGCACATGCATGGGCACCATCGTTGAGAGTGCAGCTGGGAAGAACTCTGAA
CCAGAAGTCATCAGAGCTGAGGCATGGCCTTGAACATGTCACTCAGTCTCTGGGGCTTCTGTTTCACAAATGCAT
GAGGGGGCCACCAGCCAGTGGCTTTAAACCAGGGGCAGGTTGTCCCTCCAGGCAGCATTTGAAATGTGTGTGTG
TTGAGGGGGTACAGTACTGTGGGGGCACCCCTGGCATCTAGTGGGCATCCACAATGTGCAGAACAGTCTCTG
ACAGCAAAGAATTGGTCCATTCAATGCCAATTGTAGTACCTTTGAGACATTCTGGCTGAGCCAATGCCTTCTCCC
TGTCAGAGTCCCCAGAGCAGAGAGGGTCAAGGCTTCCCTGGACCTTGGCTCCCAGAGCAAGCCAAAATAAGACT
ACACTGTTGCCTTGGGGGCTTGTCTGGGCCAGGGCCAAGACGGTCTGCGTGTGCAGGGCCAGGACAGAAATAGCC
ACACATGCCGGTGAGAACAAAGAGCCTCTTCTTTCTCATGTTGACATCGACTTCTGTGCCAAGTCCTTTGGGT
ATAAGGATGCTAGGGAATTCCTATAGGCACCAACAGAAGGAAAGCTAGGGGCTTGGACTACTGGGTATAGGACT
TGCTCTAGCTCTCAGGTCTAGCCCAAGCTCAATGCAAACACAGCCCCTCCGGGCTCTCTGTTTCTGTGAGGTTT
TGGAATCCCTTCTCTGTGTCCGTGAGTCTGACAGAATCGATGATGTTCCCTTAGAGCTGGGAAATCCATGTGTT
TATTCACGGAGGGAACACCAATTACCTCCCTTGTCTTCTTGCCTGCCTTGGAGAAATCCAGAGTCTTCGGAAT
GGCAAAGGCAGCTCCTGGATTTCCCTGGAGGGGAGGCAGTACTGAGGGAAGTAGCTCCCTTCATTTCATGATGCA
CAGTTTACGCAGCAGACACAACTGCGCCTACTATTTGCTCGGTGCCCTGCAAGGTGCTGCCTAACCTTTGATTT
GTTATTTTCTGCTCTCTCAGGATAGTGCCAAATGGTGCAATGGGAAACCTGTTTGTGCTGGGGGGCTCTAGATCAC
TGGCTCCAGAACTCCCGCTGCCAGGGTAGCCCCCTACCCCCAGCCCCCTTGCTCCTGGACAGCAGTGGGTCTCACC
TTTAGCCTCTGCCCCAGTTCTGGTCTGACCCAACAGAGGGGCTCTATGATATTAAGAAGGGGCCCTTCTGCTC
TGTGCTCAACCTATTCTCCATAATAGGGAGTCTAATCCTATTCCCTTCCCTGCCTGATGAGGATGGTGTGAGGAT
GAGGAGGACGGCATCTCATTGGGGGCTTTTGGCAGTGGGCCTCATTTTAAATCCTGCAGGGCTGCCTGCCAGTGG
ATCTATCCAGCTGCTTCCTTGTAGCCAAGAATGAGTTCAATGAATTGTGATTCACTGATTTTATTGATTTTGT
TAAACAGGGAGACTGGTATTTTTGAAGCTGCTATCATTTTCTATTCTTTTATTATTTCTTTGTAATCATCTTA
TTAAAGTTTTCTTATTTAGTGGG

1079/6881
FIGURE 1008

METVMTLEGGQKPVME SYTLLSPPETQDEGWTALSLALTYRLGPGSVLLAPRSPWRLQKPPPAGGLPHSQLPA
RQWGLLRQAGPKTVSPLGTYSPPSIC

1080/6881
FIGURE 1009

GGGACGCGAGCGGGATCCAAACTTCCGGTGCCTGCAGAGCTCGGAGCGGCGGAGGCAGAGACCGAGGCTGCACCG
GCAGAGGCTGCGGGGCGGACGCGCGGGCCGGCGCAGCCATGGTGAAGATTAGCTTCCAGCCCGCCGTGGCTGGCA
TCAAGGGCGACAAGGCTGACAAGGCGTCGGCGTCGGCCCTGCGCCGGCCCTCGGCCACCGAGATCCTGCTGACGC
CGGCTAGGGAGGAGCAGCCCCCACAACATCGATCCAAGAGGGGGGGCTCAGTGGGCGGCGTGTGCTACCTGTCTGA
TGGGCATGGTCTGTGCTCATGGGCTCGTGTTCCGCTCTGTCTACATCTACAGATACTTCTTCTTGGCGAGC
TGGCCCGAGATAACTTCTTCCGCTGTGGTGTGCTGTATGAGGACTCCCTGTCTCCAGGTCCGGACTCAGATGG
AGCTGGAAGAGGATGTGAAAATCTACCTCGACGAGAACTACGAGCGCATCAACGTGCCTGTGCCCCAGTTTGGCG
GCGGTGACCCTGCAGACATCATCCATGACTTCCAGCGGGGTCTGACTGCGTACCATGATATCTCCCTGGACAAGT
GCTATGTCATCGAACTCAACACCACCATTTGTGCTGCCCCCTCGCAACTTCTGGGAGCTCCTCATGAACGTGAAGA
GGGGGACCTACCTGCGCGAGACGTACATCATCCAGGAGGAGATGGTGGTCACGGAGCATGTGAGTACAAGGAGG
CCCTGGGGTCTTCTATCTACCACCTGTGCAACGGGAAAGACACCTACCGGCTCCGGCGCCGGGCAACGCGGAGGC
GGATCAACAAGCGTGGGGCCAAGAAGTGCATGCCATCCGCCACTTCGAGAACACCTTCGTGGTGGAGACGCTCA
TCTGCGGGGTGGTGTGAGGCCCTCCTCCCCCAGAACCCCTGCCGTGTTCTCTTTCTTCTTCCGGCTGCTCT
CTGGCCCTCCTCTTCCCCCTGCTTAGCTTGTACTTTGGACGCGTTTCTATAGAGGTGACATGTCTCTCCATTCC
TCTCCAACCCTGCCCACCTCCTGTACCAGAGCTGTGATCTCTCGGTGGGGGGCCATCTCTGCTGACCTGGGTG
TGGCGGAGGGAGAGGCGATGCTGCAAAGTGTCTTCTGTGTCCCACTGTCTTGAAGCTGGGCCTGCCAAAGCCTGG
GCCCACAGCTGCACCGGCAGCCCAAGGGGAAGGACCGGTTGGGGGAGCCGGGCATGTGAGGCCCTGGGCAAGGGG
ATGGGGCTGTGGGGGCGGGGCGGCATGGGCTTCAGAAGTATCTGCACAATTAGAAAAGTCTCAGAAGCTTTTTC
TTGGAGGGTACACTTTCTTCACTGTCCCTATTCTAGACCTGGGGCTTGAGCTGAGGATGGGACGATGTGCCCAG
GGAGGGACCCACCAGAGCACAAGAGAAGGTGGCTACCTGGGGGTGTCCAGGGACTCTGTCAGTGCCTTCAGCCC
ACCAGCAGGAGCTTGGAGTTTGGGGAGTGGGGATGAGTCCGTCAAGCACAAGTCTCTGAGTGGAAACCAAGA
AGCAAGGAGCTAGGACCCCCAGTCTGCCCCCAGGAGCACAAGCAGGGTCCCCTCAGTCAAGGCAGTGGGATGG
GCGGCTGAGGAACGGGGCAGGCAAGGTCACTGCTCAGTCACGTCCACGGGGGACGAGCCGTGGGTTCTGCTGAGT
AGGTGGAGCTCATTGCTTTCTCCAAGCTTGGAAGTGTCTTGAAGATAACACAGAGGGAAGGGAGAGCCACCTG
GTACTTGTCCACCTGCCTCCTCTGTTCTGAAATTCATCCCCCTCAGCTTAGGGGAATGCACCTTTTCCCTTT
CCTTCTCACTTTTGCATGTTTTTACTGATCATTCGATATGCTAACCGTTCTCAGCCCTGAGCCTTGGAGAGGAGG
GCTGTAACGCCTTCAGTCAGTCTCTGGGGATGAACTCTTAAATGCTTTGTATATTTTCTCAATTAGATCTCTTT
TCAGAAGTGTCTATAGAACAATAAAAATCTTTTACTTCTG

1081/6881
FIGURE 1010

MVKISFQPAVAGIKGDKADKASASAPAPASATEILLTPAREEQPPQHRSKRGGSVGGVCYLSMGMVVLLMGLVFA
SVYIYRYFFLAQLARDNFFRCGVLYEDSLSSQVRTQMELEEDVKIYLDENYERINVPVPQFGGGDPADIIHDFQR
GLTAYHDISLDKCYVIELNTTIVLPPRNFWEELLMNVKRGTYLPQTYIIQEEMVVTEHVSDKEALGSFIYHLCNGK
DTYRLRRRATRRRINKRGAKNCNAIRHFENTFVVETLICGVV

1082/6881
FIGURE 1011

GATCACCACCATCGAGGCAGTGAAGTGCAAGATCCAGGTTCTGCAGCAGCAGGCAGATGATGCAGAGGAGCGAGC
TGAGCACCTCCAGCGAGAAGTTGAGGGAGAAAGGCAGGCCCGGGAAGAGAGGTATGAAGGTTGTTGAAAATCGGG
CCTTAAAAGATGAAGAAAAGATGGAACCTCAGGAAATCTAACTCCAAGAAGCTAAGCACATTGCAGAAGAGGCAG
ATAGGAAGTATGAAGAGGTGGCTCATAAGTTGGTGATCATTGAAGGAGACTTGGAAGGCACGGAGGAACGAGCTG
AGCTGGGAGAGTCCCGTTGCCAAGAGATGGATGAGCAGATTAGACTGATGGACCAGAACCTGAAGTGTCTGAGTG
CTGCTGAAGAAATGTACTCTCAAAAAGAAGACAAATATGAGGAAGAGATCAAGATTCTTACTGATAAACTCAAGG
AGGCAGAGACCAGTGCTGAGTTTGCTGAGAGATCGGTAGCCAAGCTGGAAAAGACAATTGATGACTTGGAGATA
AACTGAAACGCACCAAAGAGGAGCACCTCTGTACACAAAGGATGCTGGACCAGACTCTGATTGACCTGAATGAGA
TGTAGAACGCCCCAGTCCCACCCTGCTGCTGCTCCTCCCTCTTCTCAGAGAATCCAGCTGGGCTAGAGGCTGAGC
ACCTTTGGAACAACATTTAAGGGAATGTGAGCACAATGCATAATGTCTTTAAAAGCATGTTGTGATG

1083/6881
FIGURE 1012

GTTTTGCCCTGCTAGCATCTCCCTGTAACCTCTCCCAATCTTGAGGAGTGATCCCTGTCCCAGCCCCTGGAAAGGGG
CAGGAACGACAAACTCAAAGTCCAGGATGTTTACCATTGACAAGAGCCATGGAAGAGGCTCTTTTTCAGCACTTCA
TGCACCAGAAGCTGGGGATCGCCTATGCCATACACAAGCCATTTCCCTTCTTTGAAGGCCTCTAGACAACCTCCA
TCATCACTAAGAGAATGTACATGGAATCTCTGGAAGCCTGTAGAAATTTGATCCCTGTATCCAGAGTGGTGCACA
ACATTCTCACCCAACTGGAGAGGACTTTTAACCTGTCTCTTCTGGTGACATTGTTTCAATCAAATTAACCTGCGTG
AATATCCCAATCTGGTGACGATTTACAGAAGCTTCAAACGTGTTGGTGCTTCTATGAACGGCAGAGCAGAGACA
CACCAATCCTACTTGAAGCCCCAACTGGCCTAGCAGAAGGAAGCTCCCTCCATACCCCACTGGCGCTGCCCCAC
CACACCCCCCTCAACCAAGCTGTTTACCCTGTGCGCCAAGAGTCAGTGAGCCTGGAACATCCTCCCAGCAAAGCG
ATGAGATCCTGAGTGAGTCGCCCAGCCCATCTGACCCTGTCTGCTCTCCCTGCACTCATCCAGGAAGGAAGAA
GCATTTCAGTGACCAATGACAAGTTAACATCCAAAATGAATGCGGAAGAAGACTCAGAAGAGATGCCAGCCTCC
TCACTAGCACTGTGCAAGTGGCCAGTGACAACCTGATCCCCCAAATAAGAGATAAAGAAGACCTCAAGAGATGC
CCCACTCTCCCTTGGGCTCTATGCCAGAGATAAGAGATAATTCTCCAGAACCAAATGACCCAGAAGAGCCCCAGG
AGGTGTCCAGCACACCTTCAGACAAGAAAGGAAAGAAAGAAAGATGTATCTGGTCAACTCCAAAAGGAGAC
ATAAGAAAAAAGCCTCCCAAGAGGGACAGCCTCATCTAGACACGGAATCCAAAAGAAGCTCAAAAGGGTGGATC
AGGTTCTCAAAAGAAAGATGACTCAACTTGTAACCTCCACGGTAGAGACAAGGGCCCCAAAGGCGAGAAGTGAAT
GTGCCGAAAAGTCGAGATCAGAGGAGATCATTGATGGCACTTCAGAAATGAATGAAGGAAAGAGTCCCAGAAGA
CGCCTAGTACACCACGAAGGGTCACACAAGGGGCAGCCTCACCTGGGCATGGCATCCAAGAGAAGCTCCAAGTGG
TGGATAAGGTGACTCAAAGGAAAGACGACTCAACCTGGAACCTCAGAGGTATGATGAGGGTCCAAAAGGCAAGAA
CTAAATGTGCCCCGAAAGTCCAGATCGAAAGAAAAAGAAAGGAGAAAGATATCTGTCAAGCTCAAAAGGAGAT
TTCAGAAAAATATTACCGAAGAGGAAAACCCAAAAGTGACACTGTGGATTTTCACTGTTCTAAGCTCCCCGTGA
CCTGTGGTGAGGCGAAAGGGATTTTATATAAGAAGAAAATGAAACACGGATCCTCAGTGAAGTGCAATCGGAATG
AGGATGGAACTTGTTAACACCAAATGAATTTGAAGTCGAAGGAAAAGGAAGGAACGCAAAGAAGTGGAAACGGA
ATATACGTTGTGAAGGAATGACCCTAGGAGAGCTGCTGAAGCGGAAAACTCGGATGAATGCGAGGTGTGCTGTC
AAGGGGGACAACCTTCTGTGCTGCGGTACTTGTCCACGAGTCTTCCATGAGGACTGTCACATCCCCCTGTGGAAG
CCAAGAGGATGCTGTGGAGTTGCACCTTCTGCAGGATGAAGAGGTCTTCAGGAAGCCAACAGTGCCATCATGTAT
CTAAGACCTGGAGAGGCAGATGCAGCCTCAGGACCAGCTGATTGAGATTACGGTGAGCCCTTTTCAAGGAAGCAA
TGTGTTTGGACCTGGTTAAGGAAAGGCTGATTACGGAATGTACACGGTGGCATGGTTTGTGCGAGACATGCGCC
TGATGTTTCGCAACCATAAAACATTTTACAAGGCTTCTGACTTTGGCCAGGTAGGACTTGACTTAGAGGCAGAAT
TTGAAAAAGATCTCAAAGACGTGCTCGGTTTTTATGAAGCCAATGACGGCGGTTTTCTGGACTCTTCTTGAACCT
GTTCTGTAAAGACTGAAGCATCCCCACCTCAGGATTCAGCTGATGGGACCCTGGCTTGACTGTTGATTGCCAGT
GAGTCTGGGATGTAATTGGCTGCCCTCAGGACCCAAACCCAGACACTTCATAGGATTATCACACCCTCCATCTTT
ATTCTTTCTTTTACCTTTAAAGTCTATATCTA

1084/6881
FIGURE 1013

MFTMTRAMEEALFQHFMHQKLGIAAYAIHKPFPFFEGLLDNSIITKRMYESLEACRNLIIPVSRVVHNILTQLERT
FNLSLLVTLFSQINLREYPNLVTIYRSFKRVGASYERQSRDTPILLEAPTGLAEGSSLHTPLALPPPQPPQPPSCS
PCAPRVSEPGTSSQQSDEILSESPSPSDPVLPLPALIQEGRSTSVTNDKLTSKMNAEEDSEEMPSLLTSTVQVAS
DNLIPQIRDKEDPQEMPHSPLGSMPEIRDNSPEPNDPEEPQEVSSTPSDKKGKKRKRRCIWSTPKRRHKKKSLPRG
TASSRHGIQKKLKRVDQVPQKKDDSTCNSTVETRAQKARTECARKSRSEEIIDGTSEMNEGKRSQKTPSTPRRVT
QGAASPGHGIQEKLVVDKVTQRKDDSTWNSEVMRVQKARTKARKSRSEKKEKKEKDICSSEKRRFQKNIHRRG
KPKSDTVDFHCSKLPVTCGEAKGILYKKMKHGSSVKCIRNEDGTWLTNPNEFEVEGKGRNAKNWKNIRCEGMTL
GELLKRKNSDECEVCCQGGQLLCCGTCPRVFHEDCHIPPEAKRMLWSCTFCRMKRSSGSQQCHHVSKTLERQMQ
PQDQLIRDYGEPPQEAMWLDLVKERLITEMYTVAVFVRDMRLMFRNHKTFYKASDFGQVGLDLEAEFEKDLKDVL
GFHEANDGGFWTLP

1085/6881
FIGURE 1014

GTTTTGCCTGCTAGCATCTCCCTGTAACCTCTCCCAATCTTGAGGAGTGATCCCTGTCCCAGCCCCTGGAAAGGGG
CAGGAACGACAAACTCAAAGTCCAGGATGTTTACCCTATGACAAGAGCCATGGAAGAGGCTCTTTTTCAGCACTTCA
TGCAACGAGAAGCTGGGGATCGCCTATGCCATACACAAGCCATTTCCCTTCTTTGAAGGCCTCCTAGACAACCTCCA
TCATCACTAAGAGAATGTACATGGAATCTCTGGAAGCCTGTAGAAATTTGATCCCTGTATCCAGAGTGGTGACACA
ACATTCTCACCCAACTGGAGAGGACTTTTAACCTGTCTCTTCTGGTGACATTGTTTCAGTCAAATTAACCTGCGTG
AATATCCCAATCTGGTGACGATTTACAGAAGCTTCAAACGTGTTGGTGCTTCCCTATGAACGGCAGAGCAGAGACA
CACCAATCCTACTTGAAGCCCCAACTGGCCTAGCAGAAGGAAGCTCCCTCCATACCCCACTGGCGCTGCCCCAC
CACACCCCCCTCAACCAAGCTGTTTACCCTGTGCGCCAAGAGTCAGTGAGCCTGGAACATCCTCCCAGCAAAGCG
ATGAGATCCTGAGTGAGTCGCCCAGCCCCTCTGACCCTGTCTGCTCTCCCTGCACTCATCCAGGAAGGAAGAA
GCATTTCAGTGACCAATGACAAGTTAACATCCAAAATGAATGCGGAAGAAGACTCAGAAGAGATGCCAGCCTCC
TCACTAGCACTGTGCAAGTGGCCAGTGACAACCTGATCCCCCAAATAAGAGATAAAGAAGACCTCAAGAGATGC
CCCCTCTCCCTTGGGCTCTATGCCAGAGATAAGAGATAATTCTCCAGAACCAAATGACCCAGAAGAGCCCCAGG
AGGTGTCCAGCACACCTTCAGACAAGAAAGGAAAGAAAAGAAAAGATGTATCTGGTCAACTCCAAAAGGAGAC
ATAAGAAAAAAGCCTCCCAAGAGGGACAGCCTCATCTAGACACGGAATCCAAAAGAAGCTCAAAAGGGTGGATC
AGGTTCTCAAAAAGAAAGATGACTCAACTTGTAACCTCACGGTAGAGACAAGGGCCCCAAAAGGCGAGAAGTGAAT
GTGCCCCGAAAGTCGAGATCAGAGGAGATCATTGATGGCACTTCAGAAATGAATGAAGGAAAGAGTCCCAGAAGA
CGCCTAGTACACCACGAAGGGTCACACAAGGGGCAGCCTCACCTGGGCATGGCATCCAAGAGAAGCTCCAAGTGG
TGGATAAGGTGACTCAAAGGAAAGACGACTCAACCTGGAACCTCAGAGGTATGATGAGGGTCCAAAAGGCAAGAA
CTAAATGTGCCCCGAAAGTCCAGATCGAAAGAAAAGAAAAGGAGAAAGATATCTGTTCAAGCTCAAAAAGGAGAT
TTCAGAAAAATATTACCCGAAGAGGAAAACCCAAAAGTGACACTGTGGATTTTCACTGTTCTAAGCTCCCCGTGA
CCTGTGGTGAGGCGAAAGGGATTTTATATAAGAAGAAAATGAAACACGGATCCTCAGTGAAGTGCATTTCGGAATG
AGGATGGAACTTGGTTAACACCAAATGAATTTGAAGTCGAAGGAAAAGGAAGGAACGCAAAGAAGTGGAAACGGA
ATATACGTTGTGAAGGAATGACCCTAGGAGAGCTGCTGAAGAGTGGACTTTTGCTCTGTCTCCAAGAATAAATC
TCAAGAGAGAGTTAAATAGCAAGTGAATTTCTACTACCCTCTCAGTCACCATGTTGCAGACTTTCCCTGTCTGGA
GGCTCACCTTAGAGCTTCTGAGTTTCCAAGCTCTGAGTCACCTCCACATTTGGGCATGGCATCTTCAAAACAATT
AATTTGCATAGTTAATTTGGGATGGGGAAGCAAATGACTCTAAATAAAAAATTAATGAAAAGCTCAAAAAAAA
AAAAA

1086/6881

FIGURE 1015

MFTMTRAMEEALFQHFHQKLGIAIAIHKPFPFFEGLLDNSIITKRMYESLEACRNLI PVSRVVHNILTQLERT
FNLSLLVTLFSQINLREYPNLVTIYRSFKRVGASIERQSRDTPILLEAPTGLAEGSSLHTPLALPPPQFPQSCS
PCAPRVSEPGTSSQQSDEILSESPSPSDPVLPLPALIQEGRSTSVTNDKLTSKMNAEEDSEEMPSLLTSTVQVAS
DNLIPQIRDKEDPQEMPHSPLGSMPEIRDNSPEPNDPEEPQEVSSTPSDKKGKKRRCIWS TPKRRHKKKSLPRG
TASSRHGIQKKLKRVDQVPQKKDDSTCNSTVETRAQKARTECARKSRSEEIIDGTSEMNEGKRSQKTPSTPRRVT
QGAASPGHGIQEKLQVVDKVTQRKDDSTWNSEVMMRVQKARTKARKSRSEKKKEKD ICSSSKRRFQKNIHRRG
KPKSDTVDFHCSKLPVTCGEAKGILYKKMKKHGSSVKCIRNEDGTWLT PNEFEVEGKGRNAKNWKRNI RCEGMTL
GELLKSGLLLCPPRINLKRELNSK

1087/6881
FIGURE 1016

GTTTTGCCTGCTAGCATCTCCCTGTAACCTCTCCCAATCTTGAGGAGTGATCCCTGTCCCAGCCCCTGGAAAGGGG
CAGGAACGACAAACTCAAAGTCCAGGATGTTTACCATGACAAGAGCCATGGAAGAGGCTCTTTTTCAGCACTTCA
TGCACCAGAAGCTGGGGATCGCCTATGCCATACACAAGCCATTTCCCTTCTTTGAAGGCCTCCTAGACAACCTCCA
TCATCACTAAGAGAATGTACATGGAATCTCTGGAAGCCTGTAGAAATTTGATCCCTGTATCCAGAGTGGTGCACA
ACATTCTCACCCAACTGGAGAGGACTTTTAACTGTCTCTTCTGGTGACATTGTTTCAGTCAAATTAACCTGCGTG
AATATCCCAATCTGGTGACGATTTACAGAAGCTTCAAACGTGTTGGTGCTTCCTATGAACGGCAGAGCAGAGACA
CACCAATCCTACTTGAAGCCTCAACTGGCCTAGCAGAAGGAGCTCCCTCCATACCCCACTGGCGCTGCCCCAC
CACAACCCCTCAACCAAGCTGTTACCCCTGTGCGCCAAGAGTCAGTGAGCCTGGAACATCCTCCCAGCAAAGCG
ATGAGATCCTGAGTGAGTCGCCCAGCCCCTGTGACCCTGTCTGCTCTCCCTGCACTCATCCAGGAAGGAAGAA
GCACTTCAGTGACCAATGACAAGTTAACATCCAAAATGAATGCGGAAGAAGACTCAGAAGAGATGCCAGCCTCC
TCACTAGCACTGTGCAAGTGCCAGTGACAACCTGATCCCCCAAATAAGAGATAAAGAAGACCTCAAGAGATGC
CCCCTCTCCCTTGGGCTCTATGCCAGAGATAAGAGATAATTCTCCAGAACCAATGACCCAGAAGAGCCCCAGG
AGGTGTCCAGCACACCTTCAGACAAGAAAGGAAAGAAAAGAAAGATGTATCTGGTCAACTCCAAAAGGAGAC
ATAAGAAAAAAGCCTCCCAAGAGGGACAGCCTCATCTAGACACGGAATCCAAAAGAAGCTCAAAAGGGTGGATC
AGGTTCTCCTCAAAAGAAAGATGACTCAACTTGTAACCTCCACGGTAGAGACAAGGGCCCAAAAGGCGAGAAGTGAAT
GTGCCGAAAGTGCAGATCAGAGGAGATCATTGATGGCACTTCAGAAATGAATGAAGGAAAGAGGTCCCAAGAAG
CGCCTAGTACACCACGAAGGGTCACACAAGGGGCAGCCTCACCTGGGCATGGCATCCAAGAGAAGCTCCAAGTGG
TGGATAAGGTGACTCAAAGGAAAGACGACTCAACCTGGAACCTCAGAGGTATGATGAGGGTCCAAAAGGCAAGAA
CTAAATGTGCCCCGAAAGTCCAGATCGAAAGAAAAGAAAAGGAGAAAGATATCTGTTCAAGCTCAAAAAGGAGAT
TTCAGAAAAATATTCACCGAAGAGGAAAACCCAAAAGTGACACTGTGGATTTTCACTGTCTAAGCTCCCCGTGA
CCTGTGGTGAGGCGAAAGGGATTTTATATAAGAAGAAAATGAAACACGGATCCTCAGTGAAGTGCATTCCGAATG
AGGATGGAACCTTGGTTAACACCAATGAATTTGAAGTGAAGGAAAAGGAAGGAACGCAAAGAAGTGGAAACGGA
ATATACGTTGTGAAGGAATGACCCTAGGAGAGCTGCTGAAGCGGAAAACTCGGATGAATGCGAGGTGTGCTGTC
AAGGGGGACAACCTTCTCTGCTGCGGTACTTGTCCACGAGTCTTCCATGAGGACTGTCACATCCCCCTGTGGAAG
CCAAGAGGATGCTGTGGAGTTGCACCTTCTGCAGGATGAAGAGGTCTTCAGGAAGCCACAGTGCCATCATGTAT
CTAAGACCTGGAGAGGCAGATGCAGCCTCAGGACCAGCTGAAATGTGAGTTCTCTCTTGAAGGCCTACTGTC
ATCCACAAAGCTCCTTTTTTACGGGCATCCATTTAATATTCGAGATTACGGTGAGCCCTTTCAGGAAGCAATGT
GGTTGGACCTGGTTAAGGAAAGGCTGATTACGGAATGTACACGGTGGCATGGTTTGTGCGAGACATGCGCCTGA
TGTTTCGCAACCATAAAACATTTTACAAGGCTTCTGACTTTGGCCAGGTAGGACTTGACTTAGAGGCAGAATTTG
AAAAAGATCTCAAAGACGTGCTCGGTTTTTATGAAGCCAATGACGGCGGTTTTCTGGACTCTTCCTTGACCCCTGTT
CTGTAAAGACTGAAGCATCCCCACCTCAGGATTACAGCTGATGGGACCCCTGGCTTGGACTGTTGATTGCCAGTGAG
TCTGGGATGTAATTGGCTGCCCTCAGGACCCAAACCCAGACACTTCATAGGATTATCACACCCCTCCATCTTTATT
CTTCTTTTTTACCTTTAAAAGTCTATATCTA

1088/6881
FIGURE 1017

MFTMTRAMEEALFQHFMHQKLGIAIAIHKPFPFFEGLLDNSIITKRMYESLEACRNLIIPVSRVVHNILTQLERT
FNLSLLVTLFSQINLREYPNLVTIYRSFKRVGASYERQSRDTPILLEAPTGLAEGSSLHTPLALPPPQPPQPPSCS
PCAPRVSEPGTSSQQSDEILSESPSPSDPVLPLPALIQEGRSTSVTNDKLTSKMNAEEDSEEMPSSLTSTVQVAS
DNLIPQIRDKEDPQEMPHSPLGSMPEIRDNSPEPNDPEEPQEVSSSTPSDKKGKKRRCIWSTPKRRHKKKSLPRG
TASSRHGIQKKLKRVDQVPQKKDDSTCNSTVETRAQKARTECARKSRSEEIIDGTSEMNEGKRSQKTPSTPRRVT
QGAASPGHGIQEKQLQVVDKVTQRKDDSTWNSEVMRVQKARTKARKSRSEKKKEKDICSSSKRRFQKNIHRRG
KPKSDTVDFHCSKLPVTCGEAKGILYKKMKHGS SVKCIRNEDGTWLT PNEFEVEGKGRNAKNWKRNIIRCEGRTL
GELLKRKNSDECEVCCQGGQLCCGTCPRVFHEDCHIPPEAKRMLWSC TFCRMKRSSGSQQCHHVSKTLERQMQ
PQDQLKCEFLLLKAYCHPQSSFFT GIPFNIRDYGEPPQEAMWLDLVKERLITEMYTVAWFVRDMRLMFRNHKTFY
KASDFGQVGLDLEAEFEKDLKDVLGFHEANDGGFWTLP

1089/6881

FIGURE 1018

ATGCTGGGTACGCTGCGCGCCATGGAGGGCGAGGACGTGGAAGACGACCAGCTGCTGCAGAAGCTCAGGGCCAGT
CGCCGCCGCTTCCAGAGGCGCATGCAGCGGCTGATAGAGAAGTACAACCAGCCCTTCGAGGACACCCCGGTGGTG
CAAATGGCCACGCTGACCTACGAGACGCCACAGGGATTGAGAATTTGGGGTGAAGACTAATAAAGGAAAGAAAC
AAAGGAGAGATCCAGGACTCCTCCATGAAGCCCGCGGACAGGACAGATGGCTCCGTGCAAGCTGCAGCCTGGGGT
CCTGAGCTTCCCTCGCACCGCACAGTCCCTGGGAGCCGATTCAAAAAGCGGTGAGGTGATGCCACGTGAGACCAG
GAAGAGTCAGTTGCTTGGGCCTTAGCACCTGCAGTGCCTCAAAGCCCTTTGAAAAATGAATTAAGAAGGAAATAC
TTGACCCAAGTGGATATACTGCTACAAAGTGCAGAGTATTTTGTGAGTGTGCAGGTAACAGAGCTGGAAGGGATGTA
CGTGTGACTCCGCTGCCTTCACTGGCCTCACCTGCCGTGCCTGCCCCGGATACTGCAGTCGTATCTCCGGAAAG
AGTCTTGGTGACCCAGCGAAACCAGCTTCATCTCCAGAGAATGGGATCCTTTGCATCCTTCTCCACAGACATG
GCCTTAGTACCTAGAAATGACAGCCTCTCCCTACAAGAGACCACTAGCAGCAGCTTCTTAAGCAGCCAGCCCTTT
GAAGATGATGACATTTGCAATGTGACCATCAGTGACCTGTACGCAGGGATGCTGCACTCCATGAGCCGGCTGTTG
AGCACAAAGCCATCAAGCATCATCTCCACCAAAACGTTTCATCATGCAAACTGGAAGTGCAGGAGGAGGCACAGA
TATAAGAGCAGGATGAACAAAACATATTGCAAAGGAGCCAGACGTTCTCAGAGGAGCTCCAAGGAGAACTTCATA
CCCTGCTCTGAGCCTGTGAAAGGGACAGGGGCATTAAGAGATTGCAAGAACGTATTAGATGTTTCTTGCCGTAAG
ACAGGTTTAAAATTGGAAAAAGCTTTTCTTGAAGTCAACAGACCCCAATCCATAAGTTAGATCCAAGTTGGAAG
GAGCGCAAAGTGACACCTCGAAGTATTTCTCTTGATTTACTTCGACTCCAGTGCAACATATAATCTTGATGAG
GAAAATAGATTTAGGACATTAAATGGTTAATTTCTCTGTAAAAATAGTTTCCAGACCAACAATACGACAGGGC
CATGGAGAGAACCCTCAGAGGGAGATTGAAATCCGATTGATCAGCTTCATCGGGAATATTGCCGTGAGTCCCAGG
AACCAGCCTCGCCGGATGTGCCTCCCGGACTCCTGGGCCATGAACATGTACAGAGGGGCTCCTGCCAGTCCCTGGT
GGCCTTCAGGGCTTAGAAACCCGAGGCTGAGTTTACCTTCCAGCAAAGCAAAGCAAAGGTTTAAAGTGAGGCT
TTTGAAAACCTAGGCAAAGATCTCTGGAAGCAGGTAGGTGCCTGCCCAAGAGCGATTTCATCTTCATCACTTCCA
AAGACCAACCCACACACAGCGCAACTCGCCCGCAGCAGACATCTGACCTTCAGTTCAGGGAAATAGTTCTGGA
ATATTTAGAAAGTCAGTGTCAACCCAGCAAACCTCTTTCAGTCCCAGATAAAGAAGTGCCAGGCAAGGAAT
CGTTACGATGAAATTAAAGAAGAAATTTGACAAGCTTCATCAAAAGTATTGCCTCAAATCTCCTGGGCAGATGACA
GTGCCTTTATGTATTGGAGTGTCTACAGATAAAGCAAGTATGGAAGTTCGATATCAAACAGAAGGCTTCTTAGGA
AAATTAAATCCAGACCCTCACTTCCAGGGTTTCCAGAAGTTGCCATCATCACCCCTGGGGTGCAGAAAAAGTCTA
CTGGGCTCAACTGCAATTGAGGCTCCTTCATCTACATGTGTGCTCGTGCCATCACGAGGGATGGCAGGAGGAC
CATCAGTTCCTTGCAAAAAGACCCAGGCTATCAGAACCCAGGGCTCCGGACGCCAGGGCAATTCCTGGGTGCC
TCAGATGGGGTGGACAACACCGTCAGACCGGAGACCAGGGCAGCTCTTCACAGCCCACTCAGAAGAGAGAGGA
GAGAACACGTCTTACAGGATGGAAGAGAAAAGTGATTTCTAGTAGAAAAATGGAACTAAAAGTGTGTAGCTA
GGTTATTTTCGGAGTGTTATTTATCTTCCCACTTGCTCTGTGTTGTATTTTGTGTTTTGATTCTTGAGAC
TGTGAGGACTTGGTTGACTTCTCTGCCCTTAAAGTAAATATTAGTGAAATTGGTTCCATCAGAGATAACCTCGAG
TTCTTGGTGTAGAAATTATGTGAATAAAGTTGCTCAATTAGAAAAA

1090/6881

FIGURE 1019

MLHMSRLLSTKPSSIISTKTFIMQNWNCRRRHRYKSRMNKTYCKGARRSQRSSKENFIPCSEPVKGTGALRDCK
NVLDVSCRKTGLKLEKAFLEVNRPQIHKLDPWKERKVTPSKYSSLIYFDSSATYNLDEENRFTLKWLISPVKI
VSRPTIRQGHGENRQREIEIRFDQLHREYCLSPRNQPRRMCLPDSWAMNMYRGGPASPGGLOGLETRRLSLPSSK
AKAKSLSEAFENLGKRSLEAGRCLPKSDSSSSLPKTNPTHSA TRPQQTSDLHVQGNSSGIFRKS VSPSKTLSVPD
KEVPGHGRNRYDEIKEEFDKLHQYCLKSPGQMTVPLCIGVSTDKASMEVRYQTEGFLGKLNPDPHFQGFQKLPS
SPLGCRKSLLGSTAIEAPSSTCVARAITRDGTRDHQFPKRPRLSEPOGSGRQGNLGSADGVDNTVRPGDQGSS
SQPNSEERGENTSYRMEEKSDFMLEKLETKSV

1091/6881
FIGURE 1020

AAGACCGTCCCGGATGGCCTCGGGGACTGCCAGTGTGTGGAGGTGAGCTCCGGGATTGCCGGCATTCCCGCTTCT
GCTGGTTGCTTTCATGCTGCAGGCTGCGGCCGTCAGCCCTCGCTCGCATTGGTGGCGCTGAGGTGCCGGGCGAGCA
AGTGACATGTCGTCGGGCCTCCGCGCCGCTGACTTCCCCCGCTGGAAGCGCCACATCTCGGAGCAACTGAGGCGC
CGGGACCGGCTGCAGAGACAGGCGTTCGAGGAGATCATCTGCAGTATAACAAATTGCTGGAAAAGTCAGATCTT
CATTCACTGTTGGCCAGAACTACAGGCTGAAAAGCATGACGTACCAAACAGGCACGAGATAAGTCCCGGACAT
GATGGCACATGGAATGACAATCAGCTACAAGAAATGGCCCAACTGAGGATTAAGCACCAAGAGGAACTGACTGAA
TTACACAAGAAACGTGGGGAGTTAGCTCAACTGGTGATTGACCTGAATAACCAAATGCAGCGGAAGGACAGGGAG
ATGCAGAAAGAGCTTGCAAGACAGCAAAAGGAACCTCTACCAGTCGAACAGGATGATGACATTGAGGTCATTGTG
GATGAAACTTCTGATCACACAGAAGAGACCTCTCTGTGCGAGCCATCAGCAGAGCAGCCACTAAGCGACTCTCG
CAGCCTGCTGGAGGCCCTTCTGGATTCTGTCACTAATATCTTTGGGAGACGCTCTGTCTCTTCCCTTCCAGTCCCC
CAGGACAATGTGGATACTCATCTGGTTCTGGTAAAGAAGTGAGGGTACCAGCTACTGCCTTGTGTGTCTTCGAT
GCACATGATGGGGAAGTCAACGCTGTGCAGTTCAGTCCAGGTTCCCGGTTACTGGCCACTGGAGGCATGGACCGC
AGGGTTAAGCTTTGGGAAGTATTTGGAGAAAAATGTGAGTTCAAGGGTTCCCTATCTGGCAGTAATGCAGGAATT
ACAAGCATTGAATTTGATAGTGCTGGATCTTACCTCTTAGCAGCTTCAAATGATTTTGCAAGCCGAATCTGGACT
GTGGATGATTATCGATTACGGGTAAGACCCAGTTAAGAAAGTTAGTGCAATCTCCAACTTCATGTGGTGTATC
AAGGCACAACTGGCAGGTGCTTAATTAGGGGACTTTGTTTTCCCAAAAATCATGCTTGATTACCCCTGCCCTTC
CTTCTCCTTGGGAAATCTGTGTTTCCACTTTATACTCTTTGTCCAAACTCAGTTTCAAATATTTGCATG
GGACCTCACATTTGCATGAAAACCTTGGAAATACTTTCATAAGGACTAAATACTTTGGTAGATAGCAATTTGG
CTTAATGGCACAGAACTTAGCAACAGCATGTGAATTGTGATTCTGTGGGCTCTAAAACCTAATTACCTAAAGTG
GGATATAGAAGTACAAATGGATGTATCATAGGGATAAGACAATTCTGAAACAAAACTCCAAGCTGAGAAAGAGG
GGACAGGTGTCAGAGCAGGGAGAAATGATTGGATGTTGAGGAAAGCTGCATTTGAACCAAACTTGCCAAGAATT
CTCTTGGCATCTGACAGAGAGACCAAACTTGGTTGTATCATTTTGTATTGGGGCAGGGGCGGAGGSCGAAGCA
TGTAACCTAATTTGCAGACATTTTTTTTCCCATAGCCTGAAGGAATCATCACATAAGCTTATTAATAACAAGC
TATTGAAAGATATAATGGAGGATGAATTTGGCATTAGTAGGCATTTTACT

1092/6881
FIGURE 1021

MSSGLRAADFPRWKRHISEQLRRRDRLQRQAFEEIILQYNKLEKSDLHSVLAQKLQAEKHDVPNRHEISPGHDG
TWNDNLQEMAQLRIKHQEELTELHKKRGELAQLVIDLNNQMQRKDREMOKELAEAAKEPLPVEQDDIEVIVDE
TSDHTEETSPVRAISRAATKRLSQPAGGLDSVTNIFGRRSVSSFPVPQDNVDTHPGSGKEVRVPATALCVFDAH
DGEVNAVQFSPGSRLLATGGMDRRVKLWEVFGKEKCEFKGSLSGSNAGITSIEFDSAGSYLLAASNDFASRIWTV
DYRLRVRS

1093/6881
FIGURE 1022A

AAGAGAGAAGGCTTCTGCCAGCTCCTGCAGCAGATGAAGAACAAGCACTCAGAGCAGCCGGAGCCCCGACATGATC
ACCATCTTCATCGGCACCTGGAACATGGGTAACGCCCCCCTCCCAAGAAGATCACGTCTGGTTTCTCTCCAAG
GGGCAGGGAAAGACGCGGGACGACTCTGCGGACTACATCCCCATGACATTTACGTGATCGGCACCCAAGAGGAC
CCCCTGAGTGAGAAGGAGTGGCTGGAGATCCTCAAACACTCCCTGCAAGAAATCACCAGTGTGACTTTTAAACA
GTGCGCCATCCACAGCTCTGGAACATCCGCATCGTGGTGCTGGCCAAGCCTGAGCACGAGAACC GGATCAGCCAC
ATCTGTACTGACAACGTGAAGACAGGCATTGCAAACACACTGGGGAACAAGGGAGCCGTGGGGGTGTCGTTTCATG
TTCAATGGAACCTCCTTAGGGTTCGTCAACAGCCACTTGACTTCAGGAAGTGAAAAGAACTCAGGCAGAACCAA
AACTATATGAACATTCTCCGGTTCCTGGCCCTGGGCGACAAGAAGCTGAGTCCCTTTAACATCACTCACC GGCTC
ACGCACCTCTTCTGGTTTGGGGATCTTAACCTACCGTGTGGATCTGCCTACCTGGGAGGCAGAAACCATCATCCAG
AAAATCAAGCAGCAGCAGTACGCAGACCTCCTGTCCCACGACCAGCTGCTCACAGAGAGGAGGGAGCAGAAGGTC
TTCTTACACTTCGAGGAGGAAGAAATCACGTTTGCCCCAACCTACCGTTTGTAGAGACTGACTCGGGACAAATAC
GCCTACACCAAGCAGAAAGCGACAGGGATGAAGTACAACCTTGCTTCCTGGTGTGACCGAGTCCCTCTGGAAGTCT
TATCCCCCTGGTGACGTGGTGTGTGTCAGTCTTATGGCAGTACCAGCGACATCATGACGAGTGACCACAGCCCTGTC
TTTGCCACATTTGAGGCAGGAGTCACTTCCCAGTTTGTCTCCAAGAACGGTCCCGGGACTGTTGACAGCCAAGGA
CAGATTGAGTTTCTCAGGTGCTATGCCACATTGAAGACCAAGTCCCAGACCAAATTCTACCTGGAGTTCCTACTCG
AGCTGCTTGGAGAGTTTGTCAAGAGTCAGGAAGGAGAAAATGAAGAAGGAAGTGAGGGGGAGCTGGTGGTGAAG
TTTGGTGAGACTCTTCCAAAGCTGAAGCCATTATCTCTGACCCTGAGTACCTGCTAGACCAGCACATCTCTCATC
AGCATCAAGTCCCTCTGACAGCGACGAATCCTATGGCGAGGGCTGCATTGCCCTTCGGTTAGAGGCCACAGAAACG
CAGCTGCCCATCTACACGCTCTCACCACCATGGGGAGTTGACAGGCCACTTCCAGGGGGAGATCAAGCTGCAG
ACCTCTCAGGGCAAGACGAGGGAGAAGCTCTATGACTTTGTGAAGACGGAGCGTGATGAATCCAGTGGGCCAAAG
ACCCTGAAGAGCCTCACCAGCCACGACCCCATGAAGCAGTGGGAAGTCACTAGCAGGGCCCCCTCCGTGCAGTGGC
TCCAGCATCACTGAAATCATCAACCCCACTACATGGGAGTGGGGCCCTTTGGGCCACCAATGCCCTGCACGTG
AAGCAGACCTTGTCCCTTGACCAGCAGCCACAGCCTGGAGCTACGACCAGCCGCCCAAGGACTCCCCGCTGGGG
CCCTGCAGGGGAGAAAGTCTCCGACACCTCCCGGCCAGCCGCCCATATACCCAAGAAGTTTTTACCCTCAACA
GCAAAACGGGGTCTCCCTCCAGGACACAGGAGTCAAGGCCCAGTGACCTGGGGAAGAACGCAGGGGACACGCTG
CCTCAGGAGGACCTGCCGCTGACGAAGCCCGAGATGTTTGAGAACCCCTGTATGGGTCCCTGAGTTCCTTCCCT
AAGCCTGCTCCAGGAAGGACCAGGAATCCCCAAAATGCCGCGGAAGGAACCCCGCCCTGCCCGGAACCCGGC
ATCTTGTGCGCCAGCATCGTGCTCACCAGCCAGGAGGCTGATCGCGGCAGGGGGCCCGCGGGGACAAGAGCCAAGGG
GCGCCCCGGCTGCGCTCCTTACGTGCTCATCTCTGCCGAGGGCAGGGCGGCCGCGGGGACAAGAGCCAAGGG
AAGCCCAAGACCCCGGTGAGTCCAGGCCCGGTGCCGCGCAAGAGGCCCATCAAGCCTTCCAGATCGGAAATC
AACCAGCAGACCCCGCCACCCGACGCCGCGGCCGCGCTGCCAGTCAAGAGCCCGCGGTGCTGCACCTCCAG
CACTCCAAGGGCCGCGACTACCGCGACAACACCGAGCTCCCGCATCACGGCAAGCACCGGCCGAGGAGGGGCCA
CCAGGGCCTCTAGGCAGGACTGCCATGCAGTGAAGCCCTCAGTGAGCTGCCACTGAGTCGGGAGCCCAAGGAAC
GGCGTGAAGCCACTGGACCTCTCCCGGGACCTCCTGCTGGCTCCTCCTGCCAGCTTCCTATGCAAGGCTTTGT
GTTTTCAGGAAAGGGCCTAGCTTCTGTGTGGCCACAGAGTTCAGTGCCTGTGAGACTTAGACCAAGTGCTGAG
GCTGGAAGAAAACGCACACCAGACGGGCAACAAACAGTCTGGGTCCCCAGCTCGCTCTTGGTACTTGGGACCCC
AGTGCCCTCGTTGAGGGCGCCATTCTGAAGAAAGGAAGTGCAGCGCCGATTTGAGGGTGGAGATATAGATAATAAT
AATATTAATAATAATAATGGCCACATGGATCGAACACTCATGATGTGCCAAGTGCTGTGCTAAGTGCTTTACGAA
CATTCTCATATCAGGATGACCTCGAGAGCTGAGGCTCTAGCCACCTAAAACCACGTGCCCAAACCCACAGTTT
AAAACGGTGTGTGTTTCGGAGGGGTGAAAGCATTAAAGAGCCAGTGCCTCCTGGAGTGAGACAAGGGCTCGGCC
TTAAGGAGCTGAAGAGTCTGGTAGCTTGTTTAGGGTACAAGAAGCCTGTTCTGTCCAGCTTCAGTGACACAAGC
TGCTTTAGCTAAAGTCCCGCGGGTTCGGCATGGCTAGGCTGAGAGCAGGGATCTACCTGGCTTCTCAGTTCTTT
GGTTGGAAGGAGCAGGAAATCAGCTCCTATTCTCAGTGGAGAGATCTGGCCTCAGCTTGGGCTAGAGATGCCAA
GGCCTGTGCCAGGTTCCCTGTGCCCTCCTCGAGGTGGGCAGCCATCACCAGCCACAGTTAAGCCAAGCCCCCAA
CATGTATTCCATCGTGCTGGTAGAAGAGTCTTTGTGTGTTGCCGAAAGCCGTGCTCTCCAGCCTGGCTGCCAG
GGAGGGTGGGCCTCTTGGTTCCAGGCTCTTGAAATAGTGCAGCCTTTTCTTCTATCTCTGTGGCTTTTACGCTCT
GCTTCTTGGTTATTAGGAGAAATAGATGGGTGATGCTTTTCTTATGTTGCTTTTTTCAACATAGCAGAATTAATG
TAGGGAGCTAAATCCAGTGGTGTGTGTAATGCAGAAGGGAATGCACCCACATTCCCATGATGGAAGTCTGCGT

1094/6881
FIGURE 1022B

AACCAATAAAATTGTCCTTTCTCACTCA

1095/6881

FIGURE 1023

MKNKHSEQPEPDMITIFIGTWNMGNAPPPKKITSWFLSKGQGKTRDDSDADYIPHDIYVIGTQEDPLSEKEWLEIL
KHSLQEIITSVTFKTVAIHTLWNIRIVVLAKPEHENRISHICTDNVKTGIANTLGNGAVGVSMFNGTSLGFVNS
HLTSGSEKKLRRNQNYMNLRFALGDKKLSFPNITHRFTHLFWFGDLNRYVDLPTWEAETIIQKIKQQQYADLL
SHDQLLTERREQVFLHFEEEEITFAPTYRFERLTRDKYAYTKQKATGMKYNLPSWCDRVLWKSYPVLHVVCQSY
GSTSDIMTSDHSPVFATFEAGVTSQFVSKNGPGTVDSQQQIEFLRCYATLTKTSQTKFYLEFHSSCLESFVKSQE
GENEEGSEGELVVKFGETLPKLPKPIISDPEYLLDQHILISIKSSDSDESYGEGCIALRLEATETQLPIYTPLTHH
GELTGHFQGEIKLQTSQGKTREKLYDFVKTERDESSGPKTLKSLTSHDPMKQWEVTSRAPPSCGSSITEIINPNY
MGVGPFPGPPMPLHVKQTLSPDQQPTAWSYDQPPKDSPLGPCRGESPPTPPGQPPISPKKFLPSTANRGLPPRTQE
SRPSDLGKNAGDTLPQEDLPLTKPEMFENPLYGSLSSFPPKAPRKDQESPKMPRKEPPPCPEPGILSPSIVLTKA
QEADRGEQPGKQVPAPRLRSFTCSSSAEGRAAGDKSQGKPKTPVSSQAPVPAKRPIKPSRSEINQQTPPTPTPR
PPLPVKSPAVLHLQHSGRDRDNTLPHHGKHRPEEGPPGFLGRTAMQ

1096/6881
FIGURE 1024

CGAGCGGACTCGACTCGGCACCGCTGTGCACCAATGCCCCGGGCCCTGTGCCGCCTCCCGCGGCGCGGCCTCTGGC
TGCTCCTGGCCCATCACCTCTTCATGACCACTGCCTGCCAGGAGGCTAACTACGGTGCCCTCCTCCGGGAGCTCT
GCCTCACCCAGTTCAGGTAGACATGGAGGCCGTCGGGGAGACGCTGTGGTGTGACTGGGGCAGGACCATCAGGA
GCTACAGGGAGCTGGCCGACTGCACCTGGCACATGGCGGAGAAGCTGGGCTGCTTCTGGCCCAATGCAGAGGTGG
ACAGGTTCTTCCTGGCAGTGCATGGCCGCTACTTCAGGAGCTGCCCCATCTCAGGCAGGGCCGTGCGGGACCCGC
CCGGCAGCATCCTCTACCCCTTCATCGTGGTCCCCATCACGGTGACCCTGCTGGTGACGGCACTGGTGGTCTGGC
AGAGCAAGCGCACTGAGGGCATTGTGTAGGCGGGGCCAGGCTGCCCCGCGGTGCACCCAGGCTGCAGGGTGAGG
CCAGGCAGGCCTGGGTAGGGGCAGCTTCTGGAGCCTTGGGACAGAGCAGGCCCAATGCCCCCTTCTTCCAGC
CAAGAAGAGCTCACAGGAGTCCAGAGTAGCCGAGGCTCTGGTATTAACTGGAAGCCCCCTGGCTGGAGGCCAC
CGCCACCCTAGGAAGGGGGCAGGGACGTGACCTTGACTTACCTCTGGAAGGGTCCCAGCCTAGACTGCTTACCC
CATAGCCACATTTGTGGATGAGTGGTTTGTGATTAAAAGGGATGTTCTTG

1097/6881
FIGURE 1025

MARALCRLPRRGLWLLLAHHLFMTTACQEANYGALLRELCLTQFQVDMEAVGETLWCDWGRTIRSYRELADCTWH
MAEKLGCWFENAEVDRFFLAVHGRYFRSCPISGRAVRDPPGSILYPFIVVPITVLLVTALVVWQSKRTEGIV

1098/6881

FIGURE 1026A

GGCACGAGGCCGCTCCCCGGCGGGCTGGCTCCTGGCCCCGGAAGCGCGAGCGTTCACTTAGCGGCGAGTGGCTCC
GTCTCCGCGGACAGAGCGCGCGCCCCCTGGCCCCGGCCGCGAGGGGCTCCCGGCGCGGTCCCCGAGCATTTCCTCG
CCGGGTGGAGCGGGCCGAGCCCCGGCAGGATGACAGCCCCGCGGCCGCTCAAAGCCGGGAGATCGACTGTTTGAG
CCCGGAAGCGCAGAAGCTGGCGGAAGCCCGCTCGCTGCAAAACGGGCGGCCCGCGGAGGCTCGCGAGATCCG
CATGAAGGAGCTGGAGCGGCAGCAGAAGGAGGAAGACAGTGAAGCGCTACTCTCGTAGATCCAGAAGAAACACATC
GGCTTCTGATGAAGACGAGCGCATGTCACTGGGTAGTCTGGAAGCCTGAGGGTAGAAGAGAGACCAGAAAAAGA
TTTTACTGAGAAGGGTCTCGTAACATGCCGGGCTGTCTGCAACCCAGCTGGCTCTCTGGGTGGGACTTCCTC
TCGGAGAGGCAGCGGAGACACCTCCTTCTCCATCGACACCGAGGCATCCATCAGGGAATCAAGGACTCTCTAGC
AGAAGTTGAAGAGAAATATAAGAAGGCTATGGTTTCCAATGCTCAGCTAGACAATGAAAAGACAACTTCATGTA
CCAGGTTGATACCTAAAAGATATGTTGCTGGAGCTTGAAGAACAGCTGGCTGAATCTAGGCGGAGTACGAAGA
GAAAAACAAAGAATTTGAAAGGGAAAAACAGCCACAGTATACTGCAATTTCACTTTGCTGAAGTCAAGGAGGC
CCTGAAGCAAAGAGAGGAAATGCTCGAGAAACATGGAATAATCCTAAATTCAGAAATAGCTACCAATGGAGAGAC
TTCCGACACCCTCAATAATGTTGGATACCAAGGTCTACCAAGATGACAAAAGAAGAGTTAAATGCCCTCAAGTC
TACAGGGGATGGGACCTAGGAAGAGCCAGTGAAGTGGAGGTGAAAAATGAAATCGTGGCGAATGTGGGGAAAAAG
AGAAATCTTGACAATACTGAGAAAGAACACACAGAGGACACAGTGAAGGACTGTGTGGACATAGAGGTATT
CCCTGCTGGTGAGAATACCGAGGACCAGAAATCCTCTGAAGACACTGCCCATTCCTAGGAACCTTAGCAGGTGC
TACCTATGAAGAACAGGTTCAAAGCCAAATTCCTTGAGAGCAGTTCTCTCCCTGAAAACACAGTACAGGTTGAGTC
AAATGAGGTTCATGGGTGACACAGATGACAGGACCAGAACTCCCTTGAGCCATCCAATGTTGGAGTGACTTAGA
TGGTGGGAACACACAGAGAATGTGGGAGAGGCAGCAGTGACTCAGGTTGAAGAGCAGGCAGGCACAGTGGCCTC
GTGTCTTTAGGGCATAGTGATGACACAGTTTATCATGATGACAAATGTATGGTAGAGGTCCCCAAGAGTTAGA
GACAAGCACAGGGCATAGTTTAGAGAAAGAATTCACCAACCAGGAAGCAGCTGAGCCCAAGGAGGTTCCAGCGCA
CAGTACAGAAGTAGGTAGGGATCACAACGAAGAAGAGGGTGAAGAAACAGGATTAGGGACGAGAAACCAATCAA
GACAGAAGTTCTGGTTCTCCAGCAGGAACAGGGGCAACTGTCAGGAAGCGACAGGTCCAAGTACAGTAGACAC
TCAAAATGAACCTTTAGATATGAAGAGCCCGATGAAGAAAAGAGTGACCAACAGGGAGAGGCATTGGACTCATC
GCAGAAGAAGACAAAGAACAAGAAAAAGAAAAACAAGAAGAAAAATCCCCAGTACCCGTAGAAACCTTAAAGA
TGTTAAAAAAGAGTTAAGCTATCAGAACACAGATTTAAGTGAATTAAGGAAGAAGAGCAGGTAAAGTCTACTGA
CAGAAAGTCAGCAGTGGAAGCCCCAAACGAGGTGACTGAAATCCAAACAGAAATTCAGCAGAGAAAGCAGTGA
AAATGTTGATTGTCCGGAGAACTCTAAATTAAGTTGGATGGAAAACCTTGACCAAGAAGGTGATGATGTACAAAC
AGCAGCTGAGGAGGTACTAGCTGATGGAGACACATTAGATTTTGAGGATGACACCGTTCAATCATCAGGCCCCGAG
GGCTGGTGGTGAAGAATTAGATGAAGGTGTTGCAAAAGATAATGCTAAAATAGATGGTGCCACTCAAAGCAGTCC
TGCAGAACCAAAGAGCGAAGACGCAGATCGCTGCACCCTGCCCGAACATGAAAGTCCCTCACAGGACATTAGTGA
TGCTGTGAAGCAGAAAGTACAGAGAGGTGTGAGATGTCAGAACATCCAAGTCAGACCGTCAGGAAAGCTTTAGA
CAGCAATAGCCTAGAGAACGATGACTTGTCCGCCACCAGGAAGAGAGCCAGGGCACTTCAATCCAGAAAGCAGAGA
AGATACCAGAGGAGGGAATGAGAAGGGCAAAAGCAAAGAAGACTGTACCATGTCTTAAGCTGAGGCAGGCGGCAG
GCGCGGTGCACAGGAAGTCTCAGTGTGAAGGGGTCTTTTCTCTCCACTGCCAATGTAAGTAGAATGTTCTAAATT
CATAGAGAGGCACTGTATGACAATTACCAGGTGCTCTACTGCTTTAAGTTATAGACTGTTACTTGTAGATTTCCA
TGTAATCATTGAGGTTATCACCCAGATTAGAAAGACATATTTGTTATCAGTGTACGTTCTAATTGAGAGCATTCC
AGTAGTATCAACAATAATGTCTACTGTTTATAGTCCACTTAATAAAAAATAGAGGCATTTACTATTTGCCTTAGG
CTGATAGGAATGTGGGTTTTCTTGACCAAAATATATCAGCATCTAATTGAAATGACCAATAGCATTCTTAGACTT
CTGTATTATGAATATAATTGATATTTAAATTAATGTCTTGTTACATATGTGTACTTTCATATTTGATTTTAAA
TGTACATTATAACCTGTATGGTATTTTATTTAAAGGAGATAAACAGCCAAATAGCAATAGGTCACTGAATGATA
AGATTTGCACCTTAGAACAATAATCATTTTTAAGGATAACAAGTAAATGTCTGAAAGCATGAGGGGCTTTATTTGC
CTTTACCTCATATGAGTCTTTGATCTTGAACCGATACTTTTGGATCTCATTGTTGATATACCTGAATTTACTTTG
TAAGAGATTTTAACTTCACTTCATGCTGATGATGATCAAATTCATTTTATAGAAAGATTTAAAGTTTTTTCTG
GAAGTGATATATGTCAAATTACATTTCTACTGCAGTATTTGAGCAGGGACAGTCATTTTTAAATGTTTTTGGC
CGGGCGTGGTGGCTCATGCCTGTAATCTCAGTACATTGGGAGGCCAAGGCAGGTGGATCACCTGAGGTCAAGAGT
TCGAGGCCAGCCTGGCCACATGGTGAAACCTGTCTCTACTAAAAATACAAAAAATTGGCCGGCGTGATGGTG
GGCGCTGTAAATCCAGCCACTCCAGAGGCTGAGGCAGGAGAATCGCTTGAACCTGCGAGGCAGAGATTGCAGTG

1099/6881

FIGURE 1026B

AGCCAAGATCAAGCCATTGTACTCCAGCCTGGACAACAAGAGCGAAACTCTGTCTA

1100/6881
FIGURE 1027

MTSPAAAQSREIDCLSPEAQKLAEARLAAKRAARAEAREIRMKELERQQKEEDSERYSRRSRRNTSASDEDERMS
VGSRGSLRVEERPEKDFTEKGSRNMPGLSAATLASLGGTSSRRGSGDTSFSIDTEASIREIKDSLAEVEEKYKKA
MVSNAQLDNEKTNFMYQVDTLKDMLEEEQLAESRRQYEEKNEFEREKHAHSILQFQFAEVKEALKQREEMLE
KHGIILNSEIATNGETSDTLNNGVYQGPTKMTKEELNALKSTGDGTLGRASEVEVKNEIVANVGKREILHNTEKE
QHTEDTVKDCVDIEVFPAGENTEDQKSSED TAPFLGTLGATYEEQVQSQILESSSLPENTVQVESNEVMGAPDD
RTRTPLEPSNCWSDLDGGNHTENVGEAAVTQVEEQAGTVASCPLGHSDDTVYHDDKCMVEVPQELETSTGHSLEK
EFTNQEAAPKEVPAHSTEVRDHNNEEGEETGLRDEKPIKTEVPGSPAGTEGNCQEATGPSTVDTQNEPLDMKE
PDEEKSDQQGEALDSSQKKTNNKKKKKKKSPVPVETLKDVKKELTYQNTDLSEIKEEEQVKSTDRKSAVEAQN
EVTENPKQKIAAESSENVDCPENPKIKLDGKLDQEGDDVQTAAEEVLADGDTLDFEDDTVQSSGPRAGGEELDEG
VAKDNAKIDGATQSSPAEPKSEDADRCTLPEHESPSQDISDACEAESTERCESMSEHPSQTVRKALDSNSLENDL
SAPGREPGHFNPESEDTRGGNEKGKSKEDCTMS

1101/6881
FIGURE 1028A

CAGTTTGGAGCTCAGTCTTCCACCAAAGGCCGTTCAAGTTCTCCTGGGCTCCAGCCTCCTGCAAGGACTGCAAGAG
TTTTCTCCGAGCTCTGAGTCTCCACTTTTTTGGTGGAGAAAGGCTGCAAAAAGAAAAGAGACGCAGTGAGTG
GGAAAAGTATGCATCCTATTCAAACCTAATTGAATCGAGGAGCCCAGGGACACACGCCTTCAGGTTTGCTCAGGG
GTTTCATATTTGGTGTCTAGACAAATTCAAATGAGGAAACATCGGCACTTGCCCTTAGTGCCGCTCTTTGCTC
TTTCTCTCAGGCTTTCTACAACCTCATGCCCAGCAGCAGCAAGCAGATGTCAAAAATGGTGCAGGCTGCTGATATA
ATATTTCTAGTGGATTCTCTTGACCATTGGAGAGGAACATTTCCAACCTGTTCGAGAGTTTCTATATGATGTT
GTAAAATCCTTAGCTGTGGGAGAAAATGATTTCATTTTGCTCTGGTCCAGTTCAACGGAAACCCACATACCGAG
TTCTGTAAATACGTATCGTACTAAACAAGAAGTCCCTTTCTCATATTTCCAACATGTCTTATATTGGGGGAACC
AATCAGACTGGAAAAGGATTAGAATACATAATGCAAAAGCCACCTCACCAGGCTGCTGGAAGCCGGGCCGGTGAC
GGAGTCCCTCAGGTTATCGTAGTGTTAACTGATGGACACTCGAAGGATGGCCTTGCTCTGCCCTCAGCGGAACTT
AAGTCTGCTGATGTTAACGTGTTTGAATTGGAGTTGAGGATGCAGATGAAGGAGCGTTAAAAGAAATAGCAAGT
GAACCGCTCAATATGCATATGTTCAACCTAGAGAAATTTACCTCACTTCATGACATAGTAGGAAACTTAGTGTC
TGTGTGCAATTCACCGTGAGTCCAGAAAGGGCTGGGGACACGGAAACCCCTTAAAGACATCACAGCACAAAGACTCT
GCTGACATTATTTTCCTTATTGATGGATCAAACAACACCGGAAGTGTCAATTTGCGAGTCATTCTCGACTTCCTT
GTAAATCTCCTTGAGAAACTCCCAATTGGAACCTCAGCAGATCCGAGTGGGGGTGGTCCAGTTTAGCGATGAGCCC
AGAACCATGTTCTCCTTGACACCTACTCCACCAAGGCCAGGTTCTGGGTGCAGTGAAAGCCCTCGGGTTTGCT
GGTGGGGAGTTGGCCAATATCGGCCCTCGCCCTTGATTTCGTGGTGGAGAACCCTTCACCCGGGCAGGGGGCAGC
CGCGTGGAGGAAGGGGTTCCCCAGGTGCTGGTCTCATAAGTGCCGGGCTTCTAGTGACGAGATTGCTACGGG
GTGGTAGCACTGAAGCAGGCTAGCGTGTTCTCATTGGCCTTGAGGCCAGGCCGCTCCAGGGCAGAGCTTCAG
CACATAGCTACCGATGACAACCTTGGTGTCTACTGTCCCGGAATTCGTTAGCTTTGGGGACCTCCAGGAGAAATTA
CTGCCGTACATTGTTGGCGTGGCCCCAAGGCACATTGTCTTGAAACCGCCAACCATTGTACACAAGTCATTGAA
GTCAACAAGAGAGACATAGTCTTCTGGTGGATGGCTCATCTGCACTGGGACTGGCCAACCTCAATGCCATCCGA
GACTTCATTGCTAAAGTCATCCAGAGGCTGGAATCGGACAGGATCTTATCCAGGTGGCAGTGCGCCAGTATGCA
GACACTGTGAGGCCGTGAATTTTATTCAATACCCATCCAACAAAAGGGAAGTCATAACCGCTGTGCGGAAAATG
AAGCCCCCTGGACGGCTCGGCCCTGTACACGGGCTCTGCTCTAGACTTTGTTGTAACAACCTATTACAGAGTTCA
GCCGGCTACCGGGCTGCGGAGGGGATTCTTAAGCTTTTGGTGTGATCACAGGTGGTAAGTCCCTAGATGAAATC
AGCCAGCCTGCCAGGAGCTGAAGAGAAGCAGCATAATGGCCTTTGCCATTGGGAACAAGGTTGCCGATCAGGCT
GAGCTGGAAGAGATCGCTTTCGACTCTCCCTGGTGTTCATCCAGCTGAGTCCGAGCCGCCCTTATGCAAGGC
ATGCTGCCTGGCTTGTGTCACCTCTCAGGACCTCTCTGGAACCCCTGAAGTTCACCTCAAACAAAAGGGATATC
ATCTTTCTTTTGGATGGATCAGCCAACGTTGGA AAAACCAATTTCCCTTATGTGCGGACTTTGTAATGAACCTA
GTTAACAGCCTTGATATTGGAATGACAATATTCGTGTTGGTTTAGTGCAATTTAGTGACACTCTGTAAACGGAG
TTCTCTTTAAACACATACCAGACCAAGTCAGATATCCTTGGTCATCTGAGGCAGCTGCAGCTCCAGGGAGGTTG
GGCCTGAACACAGGCTCAGCCCTAAGCTATGTCTATGCCAACCACTTCACGGAAGCTGGCGGCAGCAGGATCCGT
GAACACGTGCCGAGCTCCTGCTTCTGCTCAGAGCTGGGCACTCTGAGGACTCCTATTGCAAGCTGCCAACGCC
TTGACACGCGCGGCATCCTGACTTTTTGTGTGGGAGCTAGCCAGGCAATAAGGCAGAGCTTGAGCAGATTGCT
TTTAACCCAAGCCTGGTGTATCTCATGGATGATTTACGCTCCCTGCCAGCTTTGCCTCAGCAGCTGATTACGCC
CTAACCATATGTTAGTGAGGTGTGGAGGAAGTACCCTCGCTCAGCCAGAGAGCAAGCGAGACATTCTGTTT
CTCTTTGACGGCTCAGCCAATCTTGTGGCCAGTTCCCTGTTGTCGTTGACTTTCTCTACAAGATTATCGATGAG
CTCAATGTGAAGCCAGAGGGGACCCGAATTGCGGTGGCTCAGTACAGCGATGATGTCAAGGTGGAGTCCCGTTTT
GATGAGCACCAGAGTAAGCCTGAGATCCTGAATCTTGTGAAGAGAATGAAGATCAAGACGGGCAAAGCCCTCAAC
CTGGGCTACGCGCTGGACTATGCACAGAGGTACATTTTTGTGAAGTCTGCTGGCAGCCGGATCGAGGATGGAGTG
CTTCAGTTCCTGGTGTCTGCTGCTGCGAGGAAGGTCTCTGACCGTGTGGATGGGCCAGCAAGTAACCTGAAGCAG
AGTGGGGTGTGCTTTTCTATCTTCCAAGCCAAGAACGCAGACCTGCTGAGTTAGAGCAGATCGTGTCTCTCA
GCGTTTATCCTGGCTCAGAGTCCGCTTCCCAAGATTGGAGATCTTCATCCACAGATAGTGAATCTCTTAAATCA
GTGCACAACGGAGCACCAGCAGCTTTTCAAGTGAAAAGGACGTGGTGTCTGCTTGATGGCTCTGAGGGCGTC
AGGAGCGGTTTCCCTCTGTTGAAAGAGTTTGTCCAGAGAGTGGTGAAAAGCCTGGATGTGGGCCAGGACCGGGTC
CGCGTGGCCGTGGTGCAGTACAGCGACCGGACCGGCCGAGTTCTACCTGAATTCATACATGAACAAGCAGGAC
GTGCTCAACGCTGTCCGCCAGCTGACCTGCTGGGAGGGCCGACCCCCAACACGGGGCCGCCCTGGAGTTTGTC

1102/6881
FIGURE 1028B

CTGAGGAACATCCTGGTCAGCTCTGCGGGAAGCAGGATAACAGAAGGTGTGCCCCAGCTGCTGATCGTCTCACG
GCCGACAGGTCTGGGGATGATGTGCGGAACCCCTCCGTGGTCTGTAAGAGGGGTGGGGCTGTGCCCATTGGCATT
GGCATCGGGAACGCTGACATCACAGAGATGCAGACCATCTCCTTCATCCCGGACTTTGCCGTGGCCATTCCCACC
TTTCGCCAGCTGGGGACCGTCCAACAGGTCATCTCTGAGAGGGTGACCCAGCTCACCCGCGAGGAGCTGAGCAGG
CTGCAGCCGGTGTGTCAGCCTCTACCGAGCCCAGGTGTTGGTGGCAAGAGGGACGTGGTCTTTTCTCATCGATGGG
TCCCAAAGTGCCGGGCTGAGTTCAGTACGTTTCGCACCCTCATAGAGAGGCTGGTTGACTACCTGGACGTGGGC
TTTGACACCACCCGGGTGGCTGTCTCCAGTTTCAGCGATGACCCCAAGGTGGAGTTCCTGCTGAACGCCCATTC
AGCAAGGATGAAGTCGAGAACGCGGTGCAGCGGCTGAGGCCCAAGGGAGGGCGGCAGATCAACGTGGGCAATGCC
CTGGAGTACGTGTCCAGGAACATCTTCAAGAGGCCCTGGGGAGCCGATTGAAGAGGGCGTCCCGCAGTTCTCTG
GTCCTCATCTCGTCTGGAAGTCTGACGATGAGGTGGACGACCCGCGGTGGAGCTCAAGCAGTTTGGCGTGGCC
CCTTTCAGATCGCCAGGAACGCAGACCAGGAGGAGCTGGTGAAGATCTCGCTGAGCCCCGAATATGTGTCTCG
GTGAGCACCTTCGGGAGCTGCCAGCCTGGAGCAGAACTGCTGACGCCCATCACGACCTGACCTCAGAGCAG
ATCCAGAAGCTCTTAGCCAGCACTCGCTATCCACCTCCAGCAGTTGAGAGTGATGCTGCAGACATCGTCTTCTG
ATCGACAGCTCTGAGGGAGTTAGGCCAGATGGCTTTGCACATATTTCGAGATTTTGTAGCAGGATTGTTTGAAGA
CTCAACATCGGCCCCAGTAAAGTGAGAGTTGGGGTCTGTCAGTTTCAGCAATGATGTCTTCCAGAATTCTATCTG
AAAACCTACAGATCCAGGCCCGGTGCTGGACGCCATACGGCGCCTGAGGCTCAGAGGGGGTTCCCCACTGAAC
ACTGGCAAGGCTCTCGAATTTGTGGCAAGAAACCTCTTGTTAAGTCTCGGGGAGTCGCATAGAAGACGGGGTG
CCCCAACACCTGGTCTTGGTCTGGTGGAAAATCCAGGACGATGTGTCCAGGTTCCGCCAGGTGATCCGTTC
TCGGGCATTGTGAGTTTAGGGGTAGGAGACCGGAACATCGACAGAACAGAGCTGCAGACCATCACCAATGACCCC
AGACTGGTCTTACAGTGCGAGAGTTCAGAGAGCTTCCCAACATAGAAGAAAGAATCATGAACCTGTTTGGACCC
TCCCGAGCCACTCTGCACCTCCAGGGGTGGACACCCCTCCTCCTTCACGGCCAGAGAAGAAGAAAGCAGACATT
GTGTTCCTGTTGGATGGTTCCATCAACTTCAGGAGGGACAGTTTCAGGAAGTGCTTCGTTTTGTGTCTGAAATA
GTGGACACAGTTTATGAAGATGGCGACTCCATCCAAGTGGGGCTTGTCCAGTACAACCTGACCCCACTGACGAA
TTCTTCCTGAAGGACTTCTCTACCAAGAGGCAGATTATTGACGCCATCAACAAAGTGGTCTACAAAGGGGGAAGA
CACGCCAACACTAAGGTGGGCCTTGAGCACCTGCGGGTAAACCACTTTGTGCCTGAGGCAGGCAGCCGCTGGAC
CAGCGGGTCCCTCAGATTGCCTTTGTGATCACGGGAGGAAAGTCCGGTGAAGATGCACAGGATGTGAGCCTGGCC
CTCACCCAGAGGGGGGTCAAAGTGTTTGTGCTGTTGGAGTGAGGAATATCGACTCGGAGGAGGTTGGAAGATAGCG
TCCAACAGCGCCACAGCGTTCCGCGTGGGCAACGTCCAGGAGCTGTCCGAACAGGCGAGCAAGTTTGGAACT
TTGCATGATGCGATGCATGAAACCTTTGCCCTGGTGTAACTGATGCTGCCAAAGCTTGTAACTGAGTGTGATT
CTGGGGTTTATGAGTCTAGAGACCAGATGTTTTTGTGGCCAGAAAGGGCTTCGAGTCCAAGGTGGACGCCATC
TTGAACAGAATCAGCCAGATGCACAGGGTCAGCTGCAGCGGTGGCCGCTCGCCACCGTGCCTGTGTGATGGTG
GCCAACACGCCCTCGGGCCCGGTGGAGGCCCTTGAATTTGACGAGTACCAGCCAGAGATGCTCGAGAAGTTCGG
AACATGCGCAGCCAGCACCCCTACGTCTCACGGAGGACACCTGAAGGTCTACCTGAACAAGTTCAGACAGTCC
TCGCCGGACAGCGTGAAGGTGGTCATTCTTTACTGATGGAGCAGACGGAGATCTGGCTGATTTACACAGAGCA
TCTGAGAACCTCCGCCAAGAAGGAGTCCGTGCCTTGATCCTGGTGGGCCTTGAACGAGTGGTCAACTTGGAGCGG
CTAATGCATCTGGAGTTTGGGCGAGGGTTTATGTATGACAGGCCCTGAGGCTTAACCTGCTGGACTTGGATTAT
GAACTAGCGGAGCAGCTTGACAACATTGCCGAGAAAGCTTGCTGTGGGGTTCCCTGCAAGTGCTCTGGGCAGAGG
GGAGACCGCGGGCCATCGGCAGCATCGGGCCAAAGGTAATCCTGGAGAAGACGGCTACCGAGGCTATCCTGGT
GATGAGGGTGGACCCGGTGAAGCTGCTCGCTGGTGTGAACGGCACTCAAGTTTCCAGGGCTGCCCGGGCCAG
AGAGGAGTAAAGGGCTCTCGGGGATTCCAGGAGAGAAGGGCGAAGTAGGAGAAATTGGACTGGATGGTCTGGAT
GGTGAAGATGGAGACAAAGGATTACCTGGTTCTTCTGGAGAGAAAGGGAATCCTGGAAGAAGGGGTGATAAAGGA
CCTCGAGGAGAGAAAGGAGAAAGAGGAGATGTGGGATTTCAGGGGACCCGGGTAAACAGGACAAGACAGCCAG
GAGAGAGGACCCAAAGGAGAAACCGGTGACCTCGGCCCATGGGTGTCCAGGGAGAGATGGAGTACCTGGAGGA
CCTGGAGAACTGGGAAGAAATGGTGGCTTTGGCCGAAGGGGACCCCCGAGCTAAGGGCAACAAGGGCGGTCTCT
GGCCAGCCGGGCTTTGAGGGAGAGCAGGGGACAGAGGTGCACAGGGCCAGCTGGTCTCTGGTCTTCCAGGG
CTGATAGGAGAAAGGCATTCTTGACCTCGGGGAAGCGAGGTGCCGCTGGTCTCTGGAGAACGAGGCAGA
ACCGGTCCACTGGGAAGAAAGGGTGAGCCCGGAGAGCCAGGACCAAAAGGAGGAATCGGGAACCGGGGCCCTCGT
GGGAGACGGGAGATGACGGGAGAGACGGAGTTGGCAGTGAAGGACGCAGAGGCAAAAAGGAGAAAGAGGATT

1103/6881
FIGURE 1028C

CCTGGATACCCAGGACCAAAGGGTAACCCAGGTGAACCTGGGCTAAATGGAACAACAGGACCCAAAGGCATCAGA
GGCCGAAGGGGAAATTCGGGACCTCCAGGGATAGTTGGACAGAAGGGAGACCCTGGCTACCCAGGACCAGCTXXX
XXXXXXXXGGCAACAGGGGCGACTCCATCGATCAATGTGCCCTCATCCAAGCATCAAAGATAAATGCCCTTGCTGT
TACGGGCCCCCTGGAGTGGCCCCGTCTTCCCAACAGAACTAGCCTTTGCTTTAGACACCTCTGAGGGAGTCAACCAA
GACACTTTTCGGCCGGATGCGAGATGTGGTCTTGAGTATTGTGAATGACCTGACCATTGCTGAGAGCAACTGCCCA
CGGGGGGGCCCCGGGTGGCTGTGGTCACCTACAACAACGAGGTGACCACGGAGATCCGGTTTGCTGACTCCAAGAGG
AAGTCGGTCTCTGGACAAGATTAAAGAACCTTCAGGTGGCTCTGACATCCAAACAGCAGAGTCTGGAGACTGCC
ATGTCGTTTGTGGCCAGGAACACATTTAAGCGTGTGAGGAATGGATTCCCTAATGAGGAAAGTGGCTGTTTTCTTC
AGCAACACACCCACAAGAGCATCCCCACAGCTCAGAGAGGCTGTGCTCAAGCTCTCAGATGCGGGGATCACCCCC
TTGTTCTTTACAAGGCAGGAAGACCGGCAGCTCATCAACGCTTTGCAGATCAATAACACAGCAGTGGGGCATGCG
CTTGCTCTGCTGCCTGCAGGGAGAGACCTCACAGACTTCTGGAGAATGTCTTCACGTGTATGTTTGGTGGACATC
TGCAACATCGACCCATCCTGTGGATTGGCAGTTGGAGGCTTCCTTCAGGGACAGGAGAGCGGCAGGGAGCGAT
GTGGACATCGACATGGCTTTCATCTTAGACAGCGCTGAGACCACCACCCTGTTCCAGTTCAATGAGATGAAGAAG
TACATAGCGTACCTGGTCAGACAACCTGGACATGAGCCCAGATCCCAAGGCTCCAGCACTTCGCCAGAGTGGCA
GTTGTGCAGCAGCGGCCCTCTGAGTCCGTGGACAATGCCAGCATGCCACCTGTGAAGGTGGAATTCTCCCTGACT
GACTATGGCTCCAAGGAGAAGCTGGTGGACTTCCTCAGCAGGGGAATGACACAGTTGCAGGGAACCCAGGGCCTTA
GGCAGTGCCATTGAATACACCATAGAGAATGTCTTTGAAAGTGCCCCAAACCCACGGGACCTGAAAATTGTGGTC
CTGATGCTGACGGGCGAGGTGCCGGAGCAGCAGCTGGAGGAGGCCAGAGAGTCATCTGCAGGCCAAATGCAAG
GGCTACTTCTTCGTGGTCTGGGCATTGGCAGGAAGGTGAACATCAAGGAGGTATACACCTTCGCCAGTGAGCCA
AACGACGTCTTCTTCAAATTAGTGGACAAGTCCACCGAGCTCAACGAGGAGCCTTTGATGCGCTTCGGGAGGCTG
TTGCCATCCTTCGTGAGCAGTGAAAATGCTTTTACTTGTCCCCAGATATCAGGAAACAGTGTGATTGGTTCCAA
GGGGACCAACCCACAAGAACCTTGTGAAGTTTGGTCACAAACAAGTAAATGTTCCGAATAACGTTACTTCAAGT
CCTACATCCAACCCAGTGACGACAACGAAGCCGGTGACTACGACGAAGCCGGTGACCACCACAACAAAGCCTGTA
ACCACCACAACAAAGCCTGTGACTATTATAAATCAGCCATCTGTGAAGCCAGCCGCTGCAAGCCGGCCCCCTGCG
AAACCTGTGGCTGCCAAGCCTGTGGCCACAAGACGGCCACTGTAGACCCCAAGTGGCGGTGAAGCCAGCAACG
GCAGCGAAGCCTGTAGCAGCAAAGCCAGCAGCTGTAAGACCCCCGCTGCTGCTGCAAAACAGTGGCGACCAAG
CCTGAGGTCCCTAGGCCACAGGCAGCCAAACCAGCTGCCACCAAGCCAGCCACCACTAAGCCCATGGTTAAGATG
TCCCGTGAAGTCCAGGTGTTTGTAGATAACAGAGAACAGCGCCAACTCCACTGGGAGAGGCCTGAGCCCCCGGT
CCTTATTTTTATGACCTCACCGTCACCTCAGCCCATGATCAGTCCCTGGTTCTGAAGCAGAACCTCACGGTCACG
GACCGCGTCATTGGAGGCCCTGCTCGCTGGGCAGACATACCATGTGGCTGTGGTCTGCTACCTGAGGTCTCAGGTC
AGAGCCACCTACCATGGAAGTTTCAGTACAAAGAAATCTCAGCCCCACCTCCACAGCCAGCAAGGTCAGCTTCT
AGTTCAACCATCAATCTAATGGTGAGCACAGAACCATTGGCTCTCACTGAAACAGATATATGCAAGTTGCCGAAA
GACGAAGGAACCTGCAGGGATTTCATATTTAAATGGTACTATGATCCAAACACCAAAAGCTGTGCAAGATTCTGG
TATGGAGGTTGTGGTGAAACGAAAACAAATTTGGATCACAGAAAGAATGTGAAAAGGTTTTCGCTCCTGTGCTC
GCCAAACCCGAGTCATCAGTGTGATGGGAACCTAAGCGTGGGTGGCCAAACATCATATACCTCTTGAAGAAGAAG
GAGTCAGCCATCGCCAACTTGTCTCTGTAGAAGCTCCGGGTGTAGATTCCCTTGCACTGTATCATTTTCATGCTTT
GATTTACACTCGAACTCGGGAGGGAACATCCTGCTGCATGACCTATCAGTATGGTGCTAATGTGTCTGTGGACCC
TCGCTCTCTGTCTCCAGGCAGTTCTCTCGAATACTTTGAATGTTGTGTAACAGTTAGCCACTGCTGGTGTATTATG
TGAACATTCCTATCAATCCAAATTCCTCTGGAGTTTCATGTTATGCCTGTTGCAGGCAAATGTAAAGTCTAGAA
AATAATGCAAAATGTACGGCTACTCTATATACCTTTGCTTGGTTCAATTTTTTTTCCCTTTTAGTTAAGCATGACT
TTAGATGGGAAGCCTGTGTATCGTGGAGAAACAAGAGACCAACTTTTTCATTCCCTGCCCCCAATTTCCAGACT
AGATTTCAAGCTAATTTTTCTTTTCTGAAGCCTCAACAAATGATCTAGTTTCAAGGAAGCAAAATCCCTTAAT
CTATGTGACCGTTGGGACCAATGCCTTAATTAAGAATTTAAAAAGTTGTAATAGAGAATATTTTTGGCATTC
CTCTAATGTTGTGTGTTTTTTTTTTTGTGTGTGGAGGGAGGGATTAAATTTAATTTTAAATGTTTAGGAA
ATTTATACAAAGAACTTTTTAATAAGTATATTGAAAGTTT

1104/6881
FIGURE 1029

AGCCCAGCGAGGCCTGTGCTGTGAGGGGCTGTGCTGTGCCAGGGGAGGCCCCGCACCCGGGAAGCCCAGAGGAGA
GTGTGCACTCACGGAGCCCTCCTGGCAGGGCCGCGTCTCTGGAGCAGCAGACCCGTGTCCCTCTGGGAGGTCAAC
TGCGGCCTCACAGCTGGGCCTGTCCCAGCACAGCCCTTGTTGGAGGGCGCACCCCGTGTCTCTCAGCACCCAGT
GTTGAGTGCCCTTGACCTGCAGAGGGGAGCCTGTGCTGGTCCCGGGCGTCTGTGTGGCTCCTGGGTGCCCATCTGC
ACCCCTGGCCCTGTCTGGGCTTCAGTGGAGCCGTGCCACCGTGGGAGTGTCCAGGGTCCTGACGTAGCCCTGAG
ACGTGGCCAGCTCCCTGGCCTGTGTCGCCGACAGTGGGCCCTTGCTCTCTGCTCTGGGTGAGGCGAGGCAGCCACC
CATGATGAGCTCTGCAGCCTTCCCAAGGTGGCTGAGCATGGGGGTCCCTCGTACCCCTTCACGGACAGTGCTCTT
CGAGCGGGAGAGGACGGGCCTGACCTACCGCGTGCCTCGTTGCTCCCCGTGCCCCCGGGGCCACCCCTGCTGGC
CTTTGTGGAGCAGCGGCTCAGCCCTGACGACTCCACGCCCACCGCCTGGTGCTGAGGAGGGGCACGCTGGCCGG
GGGCTCCGTGCGGTGGGGTGCCCTGCACGTGCTGGGGACAGCAGCCCTGGCGGAGCACCGGTCCATGAACCCCTG
CCCTGTGCACGATGCTGGCACGGGCACCGTCTTCCTCTTCTTCATCGCGGTGCTGGGCCACACGCTGAGGCCGT
GCAGATCGCCACGGGAAGGAACGCCGCGCGCCTCTGCTGTGTGGCCAGCCGTGACGCCGGCCTCTCGTGGGGCAG
CGCCCGGGACCTCACCGAGGAGGCCATCGGTGGTGCCGTGCAGGACTGGGCCACATTCGCTGTGGGTCCCGGCCA
CGGCGTGCAGCTGCCCTCAGGCCGCTGCTGGTACCCGCTACACCTACCGCGTGGACCGCCGAGAGTGTTTGG
CAAGATCTGCCGGACCAGCCCTCACTCCTTCGCCTTCTACAGCGATGACCACGGCCGCACCTGGCGCTGTGGAGG
CCTCGTGCCCAACCTGCGCTCAGGCGAGTGCCAGCTGGCGGCGGTGGACGGTGGGCAGGCCGGCAGCTTCCTCTA
CTGCAATGCCCGGAGCCCACTGGGCAGCCGTGTGCAGGCGCTCAGCACTGACGAGGGCACCTCCTTCCTGCCCGC
AGAGCGCGTGGCTTCCCTGCCCGAGACTGCCTGGGGCTGCCAGGGCAGCATCGTGGGCTTCCCAGCCCCGCCCC

1105/6881
FIGURE 1030

GAATTGGCAGCCAACATGGCGGCGGAACGCGGCGCGGGGCAGCAACAGTCGCAGGAGATGATGGAGGTTGACAGG
CGGGTCGAGTCTGAAGAATCCGGCGATGAAGAAGGGAAGAAACACAGCAGTGGCATCGTGGCCGACCTCAGTGAA
CAGAGCCTGAAGGATGGGGAGGAGCGGGGGGAGGAGGCCAGAGAAGAACATGAGCTGCCTGTGGACATGGAA
ACCATCAACCTGGACAGAGATGCAGAGGATGTTGATTTGAATCACTATCGCATAGGGAAGATTGAAGGATTTGAG
GTACTGAAGAAAGTGAAGACTCTCTGCCTCCGCCAAAATTTAATTAAATGCATTGAGAATCTGGAGGAGCTACAG
AGTCTTCGAGAGCTGGATCTTTACGACAACCAGATCAAGAAGATTGAGAATCTGGAGGCGCTAACAGAGCTGGAG
ATTCTAGATATTTCTTTTAACTGCTGAGAAACATCGAAGGGGTTGACAAGTTGACACGACTGAAAAAACTCTTC
TTGGTCAACAATAAAATCAGTAAAATTGAGAACTTAAGCAACTTACATCAACTACAGATGCTAGAGCTGGGATCT
AACCGCATCCGGGCAATCGAAAATATCGACACCTTAACCAACCTGGAGAGTTTGTTTTTGGGGAAAAACAAAATT
ACTAAACTTCAGAACCTGGATGCGCTCACCAACCTGACAGTCCTCAGTATGCAGAGCAACCGGCTGACCAAGATC
GAGGGTCTGCAGAACCTGGTGAACCTGCGGGAGCTGTACCTTAGCCACAATGGCATCGAGGTCATCGAGGGCCTG
GAGAACAATAACAACTCACGATGTTGGACATTGCATCAAATAGAATCAAAAAGATTGAAAATATCAGCCATCTA
ACAGAGCTGCAAGAGTTCTGGATGAACGACAATCTCCTTGAGAGCTGGAGCGACCTCGACGAGCTGAAGGGAGCC
AGGAGCCTGGAGACAGTGTACCTGGAGCGGAACCCCTTGAGAAAGGACCCCCAGTACCGGCGGAAGGTCATGCTC
GCCCTCCCTCCGTCGCGGCAGATCGATGCCACGTTTCGTCAAGTTCTGAGTCCTTCTTGGCTCCTCATGTGGTCCC
TCTCCTCGGAAGAACTGCCCAGCCACGGGTTTTTAACCCACCTGTTGCTCCTGAGGTCGTCACTATATCAACAGT
CACAAACCCAATGGCAATAAAGGCACTGACGATAGCTGGCGCGCGGACGCCACACACCATTTTCAGATGCCGTT
GCAATTAATCTTGCCACACTGTC

1106/6881
FIGURE 1031

MAAERGAGQQSQEMMEVDRRVESEESGDEEGKKHSSGIVADLSEQSLKDGEERGEEDPEEEHELPVDMETINLD
RDAEDVDLNHYRIGKIEGFVLKKVKTLCRLQNLKCIENLEELQSLRELDLYDNQIKKIELEALTELEILDIS
FNLLRNIEGVDKLTRKKLFLVNNKISKIENLSNLHQLQMLELGSNRIRAIENIDTLTNLESFLGKNKITKLQN
LDALTNLTVLSMQSNRLTKIEGLQNLVNLRELYLSHNGIEVIEGLENNNKLTMLDIASNRIKKIENISHLTELQE
FWMNDNLLESWSDLDELKGARSLETVYLERNPLQKDPQYRRKVMLALPSVRQIDATFVRF

1107/6881
FIGURE 1032A

CAAAAGCTGGCAGGCTGACAGAGCGGCCTCAGGACGGACCTTCTGGCTACTGACCGTTTTGCTGTGGTTTTCCC
GGATTGTGTGTAGGTGTGAGATCAACCATGAGTTCCGTTGCAGTTTTGACCCAAGAGAGTTTTGCTGAACACCGA
AGTGGGCTGGTTCGCAACAAATCAAAGTTGCCACTCTAAATTCAGAAGAGGAGAGCGACCCCTCCAACCTACAAG
GATGCCTTCCCTCCACTTCTCTGAGAAAAGCTGCTTGCTGGAAAGTGCCAGGAACCCCTCTGGAGCCTGGGGGAAC
AAGATCCGACCCATCAAGGCTTCTGTCACTCACTCAGGTGTTCCATGTACCCCTGGAGGAGAGAAAATACAAGGAT
ATGAACCAGTTTGGAGAAGGTGAACAAGCAAAAATCTGCCTTGAGATCATGCAGAGAACTGGTGCTCACTTGGAG
CTGTCTTTGGCCAAAGACCAAGGCTCTCCATCATGGTGTGAGGAAAGCTGGATGCTGTCAAGAAAGCTCGGAAG
GACATTGTTGTAGACTGCAGACTCAGGCCTCAGCAACTGTTGCCATTCCCAAAGAACACCATCGCTTTGTTATT
GGCAAAAATGGAGAGAACTGCAAGACTTGGAGCTAAAACTGCAACCAAAAATCCAGATCCCACGCCAGATGAC
CCCAGCAATCAGATCAAGATCACTGGCACCAAGAGGGCATCGAGAAAGCTCGCCATGAAGTCTTACTCATCTCT
GCCGAGCAGGACAAACGTGCTGTGGAGAGGCTAGAAGTAGAAAAGGCATTCCACCCCTTCATCGCTGGGCCGTAT
AATAGACTGGTTGGCGAGATCATGCAGGAGACAGGCACGCGCATCAACATCCCCCACCAGCGTGAACCGGACA
GAGATTGTCTTCACTGGAGAGAAGGAACAGTTGGCTCAGGCTGTGGCTCGCATCAAGAAGATTTATGAGGAGAAG
AAAAAGAAGACTACAACCATTTGCAGTGGAAAGTGAAGAAATCCCAACACAAGTATGTCATTGGGCCCAAGGGCAAT
TCATTGCAGGAGATCCTTGAGAGAACTGGAGTTTCCGTTGAGATCCCACCCTCAGACAGCATCTCTGAGACTGTA
ATACTTCGAGGCGAACCTGAAAAGTTAGGTTCAGGCGTTGACTGAAGTCTATGCCAAGGCCAATAGCTTCACCGTC
TCCCTGTGCGCCGCCCTTCCCTGGCTTCACCGTTTCATCATTGGCAAGAAAGGGCAGAACCTGGCCAAAATCACT
CAGCAGATGCCAAAGGTTACATCGAGTTCACAGAGGGCGAAGACAAGATCACCCCTGGAGGGCCCTACAGAGGAT
GTCAATGTGGCCAGGAACAGATAGAAGGCATGGTCAAAGATTTGATTAACCGGATGGACTATGTGGAGATCAAC
ATCGACCACAAGTTCCACAGGCACCTCATTGGGAAGAGCGGTGCCAACATAAACAGAATCAAAGACCAGTACAAG
GTGTCCGTGCGCATCCCTCCTGACAGTGAGAAGAGCAATTTGATCCGCATCGAGGGGGACCCACAGGGCGTGCAG
CAGGCCAAGCGAGAGCTGCTGGAGCTTGCACTCTCGCATGGAAAATGAGCGTACCAAGGATCTAATCATTGAGCAA
AGATTTATCGCACAATCATTGGGCAGAAGGGTGAACGGATCCGTGAAATTCGTGACAAATCCCAGAGGTCATC
ATTAACCTTCCAGACCCAGCACAAAAAGTGACATTGTCCAGCTCAGAGGACCTAAGAATGAGGTGGAAAAATGC
ACAAAATACATGCAGAAGATGGTGGCAGATCTGGTGGAAAATAGCTATTCAATTTCTGTTCCGATCTTCAAACAG
TTTCACAAGAATATCATTGGGAAGGAGGGCGAAACATTAAAAAGATTTCGTGAAGAAAGCAACACCAAAATCGAC
CTTCAGCAGAGAATAGCAATTCAGAGACCATATCATCACAGGCAAGCGAGCCAACTGCCAAGCTGCCGGGAGC
AGGATTCTGTCTATTAGAAAAGACCTGGCCAACATAGCCGAGGTAGAGGTCTCCATCCCTGCCAAGCTGCACAAC
TCCCTCATTGGCACCAAGGGCGCTCTGATCCGCTCCATCATGGAGGAGTGGCGGGGTCCACATTCACTTTCCC
GTGGAAGGTTCAAGGAAGCGACACCGTTGTTATCAGGGGCCCTTCTCGGATGTGGAGAAGGCCAAGAAGCAGCTC
CTGCATCTGGCGGAGGAGAAGCAACCAAGAGTTTCACTGTTGACATCCGCGCCAAGCCAGAATACCACAAATTC
CTCATCGCAAGGGGGGGCGGCAAAATTCGCAAGGTGCGCGACAGCACTGGAGCACGTGTCACTTCCCTGCGGCT
GAGGACAAGGACCAGGACCTGATCACCATCATTGGAAGGAGGACGCCGTCCGAGAGGCACAGAAGGAGCTGGAG
GCCTTGATCCAAAACCTGGATAATGTGGTGAAGACTCCATGCTGGTGGACCCCAAGCACCACCGCCACTTCGTC
ATCCGCAGAGGCCAGGTCTTGCGGGAGATTGCTGAAGAGTATGGCGGGGTGATGGTCAGCTTCCCACGCTCTGGC
ACACAGAGCGACAAAGTCACCCTCAAGGGCGCAAGGACTGTGTGGAGGCAGCCAAGAAACGCATTCAAGGATC
ATTGAGGACCTGGAAGCTCAGGTGACATTAGAATGTGCTATACCCAGAAAATCCATCGATCTGTCTGGGCCCC
AAAGGTTCCAGAATCCAGCAGATTACTCGGGATTTCACTGTTCAAATTAATTTCCAGACAGAGAGGAGAACGCA
GTTACAGTACAGAGCCAGTTGTCCAGGAGAATGGGGACGAAGCTGGGGAGGGGAGAGGCTAAAGATTGTGAC
CCCGGCTCTCCAAGGAGGTGTGACATCATCATCTCTGGCCGAAAGAAAAGTGTGAGGCTGCCAAGGAAGCT
CTGGAGGCATTGGTTCTGTCAACATTGAAGTAGAGGTGCCCTTTGACCTTACCGTTACGTTATTGGGCAGAAA
GGAAGTGGGATCCGAAGATGATGGATGAGTTTGAGGTGAACATACATGTCCCGGCACCTGAGCTGAGCTGTGAC
ATCATCGCCATCAGGGCCTCGCTGCAAAATTTGACCGGGCCAAAGGCTGGACTGCTGGAGCGTGTGAAGGAGCTA
CAGGCCGAGCAGGAGGACCGGGCTTTAAGGAGTTTTAAGCTGAGTGTCACTGTAGACCCCAATACCATCCCAAG
ATTATCGGGAGAAAAGGGGGCAGTAATTACCAAAATCCGTTTGGAGCATGACGTGAACATCCAGTTTCTGTATAAG
GACGATGGGAACCAAGCCAGGACCAATTACCATCAGGGTACGAAAAGAACACAGAAGCTGCCAGGGATGCT
ATACTGAGAATTGTGGGTGAAGTTGAGCAGATGGTTTCTGAGGACGTCCCGCTGGACCACCGGCTTACGCCCCG
ATCATTGGTGGCCGCGCAAGCCATTGCAAAAATCATGGACGAATTCAAGGTGGACATTGCTTCCCACAGAGC

1108/6881

FIGURE 1032B

GGAGCCCCAGACCCCAACTGCGTCACTGTGACGGGGCTCCCAGAGAATGTGGAGGAAGCCATCGACCACATCCTC
AATCTGGAGGAGGAATACCTAGCTGACGTGGTGGACAGTGAGGCGCTGCAGGTATACATGAAACCCCAAGCACAC
GAAGAGGCCAAGGCACCTTCCAGAGGCTTTGTGGTGCGGGACGCACCCTGGACCGCCAGCAGCAGTGAGAAGGCT
CCTGACATGAGCAGCTCTGAGGAATTTCCCAGCTTTGGGGCTCAGGTGGCTCCCAAGACCCTCCCTTGGGGCCCC
AAACGATTAATGATCAAAAAGAACAGAACCCCTCTCCAGCCTGCTGACCCAAACCAACCACACAATGGTTTGTCTC
AATCTGACCCAGCGGCTGGACCCTCCGTAAATTGTTGACGCTCTTCCCCCTTCCCGAGGTCCCGCAGGGAGCCTA
GCGCCTGGCTGTGTGTGCGGCCGCTCCTCCAGGCCGTGGCCGTGCCCGCTCAGGACCTGCTCCACTGTTTAACACT
AAACCAAGGTCATGAGCATTTCGTGCTAAGATAACAGACTCCAGCTCCTGGTCCACCCGGCATGTCAGTCAGCACT
CTGGCCTTCATCACGAGAGCTCCGCAGCCGTGGCTAGGATTCCACTTCCTGTGTCATGACCTCAGGAAATAAACG
TCCTTGACTTTATAAAAGCC